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m
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M
    <223> exon 21
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13032-13104;

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<220>
<221> misc feature
<222> (23045)..(26452)
<223> exon 19: 23045-23154; exon 20: 23795-23895; exon
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      signal: 26447-26452
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<221> variation
<222> (826)..(23879)
<223> s at positions 826 and 23180 is G or C; y at
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      is C or T; n at position 13128 is t or tgat; r at
      positions 22211 and 23879 is A or G.
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agc cgg gac tcg ggc gcc gcg ctc tac gtc ttc tcc gag ttc aac Ser Arg Asp Ser Gly Ala Ala Leu Tyr Val Phe Ser Glu Phe Asn 70 75 80	293
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Ser Gly Cys Ser Gly Gly Pro Asn Thr Val Tyr Leu Gln Val Val Ala 50 55 60	

Al 6	a Gl 5	y Se	r Ar	g As _l	o Se:	r Gl	y Ala	a Ala	a Lei	Tyr 75	val	l Phe	e Sei	c Glu	Phe 80	
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His	Leu	Arg 35	Thr	Arg	Glu	Lys	Arg 40	Gly	Pro	Gly	Pro	Gly 45	Gly	Pro	Asn	
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<223> This can be any amino acid residue.
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<222> (466)..(983)
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taacttgtag ttttagccaa gttattaaaa ccttactgtg gatatgtgtg gaatactatg 180
agagaccaag aatccagact gttctaaata accaaaaagt aataatagag ataaatatta 240
caggaatatg tttttggtcc agtgatatga aataatcccc agatgatctt tctgttgcag 300
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cccggggtgc ctctgctgtg gtccttcggt gtgaaggcga gtgctggctc tttgactgtg 420
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 gaacttttaa attcaattta ctattttaa tgtaaattgt taggcttgtt tcaaatagct 240
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taatatcaat caacactgaa attttaaaaa tgtataaatc cagttttcca caagtagtaa 180
aacatttata acaattatgg atgccttttc cattagctat ttgcaatgct gttaaaatag 240
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Phe Leu 465 Glu Ile Cga aat gtc agt tcc aca Met Glu Ile Cga aat gtc agt tcc aca Met Glu Ile Cga aat gtc agt tcc aca Lys Ser Val Soo Cgt at Tyr Gly Gln Gln Ile Asp Asp Soo Cgt Cat Tyr Gly Gln Gln Ile Asp Soo Arg Soo Cgt ttg Gly Gln Gln Ile Asp Soo Arg Soo Cgt ttg Cat Leu Cat Gln Asp Soo Arg Soo Cgt ttg Cat Cat Gln Arg Soo Arg Soo Cgt Leu Cat Gln Arg Glu Gln Soo Cgt Leu Cat Gln Arg Soo Arg Soo Cgt Cat Cat Gln Arg Glu Gln Soo Cgt Cat Cat Cat Gln Arg Glu Gln Soo Cgt Cat Cat Cat Cat Cat Glu Cat	Glu Tyr Arg Lys Asn Val Gln Glu Asn Product agc cag tat cct gaa att gtc ttc ctg ggt Act Glu Ile Cga aatt gtc ctc ctc agt tcc agt tcc agt tcc agt tcc agt tcc agt tcc agt dcc dcc dcg gat tgt gaa gga ggg gaa gaa gaa gaa <th>Glu Tyr Arg Lys Asn Val Gln Glu Asn Pro Ala agc cag tat ctt gau att gtt ttc ctg ggt Arg att gag att ctg gaat gtc ttc ctg gtc ggg gga gtc gac gac</th> <th>Ğlu Tyr Arg Lys Asn Val Glu Asn Fro Ala Pro 460 ase Asn Val Glu Asn Fro 460 ase Asn Yal Etc Ctg ggt Asn Yal Etc Ctg Glu Ile Val Etc Ctg Glu Ile Val Leu Glu Ile Val Leu Asn Val Ser Thr Leu Asn Asn<th>Glu Tyr Arg Lys Asn Val Glu Glu Asn Pro Ala Pro Ala 460 Asn Pro Ala 460 Asn Asn Pro Ala 460 Asn Asn Asn Pro Asn Yal Etc Ctg gga acg ggg ggg Ctg Ser Asn Yal Asn Asn Asn Yal Asn Asn Yal Asn Leu Leu Asn Asn Asn Leu Asn Leu Asn Leu Asn Leu Asn Leu Asn Asn<th>Glu Tyr Arg Lys Asn Val Gln Glu Asn Pro Ala Pro Ala glu Asn Pro Ala Pro Ala Glu Ala Asn Val Fhe Leu Gly Thr Gly Ser Ala Asn Val Fhe Leu Gly Fhr Gly Ser Ala Asn Val Ser Asn Val Ser Ser Thr Leu Asn Leu Ser Ser Thr Leu Asn Leu Ser Asn Val Leu Ser Asn Leu Leu Ser Leu Leu<th>Glù Tyr Arg Lys Asn Val Gln Glu Asn Pro Ala Glu Lys 465 Asn Yal Glu Glu Glu Glu Glu He Ctt Ctt Gu att Gtt Gu att Asn Val Fee Leu Glu Thr Glu He Ctt Asn July Ser Gu Asn Leu Asn Leu Asn Val Fer Thr Leu Gu Asn Leu Ser Pro 495 aag tca gtg ctc ctg gat ttg ga gac ctt ttg Gu Asn Leu Ser Pro 4495 Asn Leu Leu</th><th>agc cag tat cct gas att gtc ttc ctg ggt acg ggg tct gcc atc cag atg ggg atg gtc ggc atg ctg gar atg ggg atg gag atg gg atg ggg atg at</th></th></th></th>	Glu Tyr Arg Lys Asn Val Gln Glu Asn Pro Ala agc cag tat ctt gau att gtt ttc ctg ggt Arg att gag att ctg gaat gtc ttc ctg gtc ggg gga gtc gac gac	Ğlu Tyr Arg Lys Asn Val Glu Asn Fro Ala Pro 460 ase Asn Val Glu Asn Fro 460 ase Asn Yal Etc Ctg ggt Asn Yal Etc Ctg Glu Ile Val Etc Ctg Glu Ile Val Leu Glu Ile Val Leu Asn Val Ser Thr Leu Asn Asn <th>Glu Tyr Arg Lys Asn Val Glu Glu Asn Pro Ala Pro Ala 460 Asn Pro Ala 460 Asn Asn Pro Ala 460 Asn Asn Asn Pro Asn Yal Etc Ctg gga acg ggg ggg Ctg Ser Asn Yal Asn Asn Asn Yal Asn Asn Yal Asn Leu Leu Asn Asn Asn Leu Asn Leu Asn Leu Asn Leu Asn Leu Asn Asn<th>Glu Tyr Arg Lys Asn Val Gln Glu Asn Pro Ala Pro Ala glu Asn Pro Ala Pro Ala Glu Ala Asn Val Fhe Leu Gly Thr Gly Ser Ala Asn Val Fhe Leu Gly Fhr Gly Ser Ala Asn Val Ser Asn Val Ser Ser Thr Leu Asn Leu Ser Ser Thr Leu Asn Leu Ser Asn Val Leu Ser Asn Leu Leu Ser Leu Leu<th>Glù Tyr Arg Lys Asn Val Gln Glu Asn Pro Ala Glu Lys 465 Asn Yal Glu Glu Glu Glu Glu He Ctt Ctt Gu att Gtt Gu att Asn Val Fee Leu Glu Thr Glu He Ctt Asn July Ser Gu Asn Leu Asn Leu Asn Val Fer Thr Leu Gu Asn Leu Ser Pro 495 aag tca gtg ctc ctg gat ttg ga gac ctt ttg Gu Asn Leu Ser Pro 4495 Asn Leu Leu</th><th>agc cag tat cct gas att gtc ttc ctg ggt acg ggg tct gcc atc cag atg ggg atg gtc ggc atg ctg gar atg ggg atg gag atg gg atg ggg atg at</th></th></th>	Glu Tyr Arg Lys Asn Val Glu Glu Asn Pro Ala Pro Ala 460 Asn Pro Ala 460 Asn Asn Pro Ala 460 Asn Asn Asn Pro Asn Yal Etc Ctg gga acg ggg ggg Ctg Ser Asn Yal Asn Asn Asn Yal Asn Asn Yal Asn Leu Leu Asn Asn Asn Leu Asn Leu Asn Leu Asn Leu Asn Leu Asn Asn <th>Glu Tyr Arg Lys Asn Val Gln Glu Asn Pro Ala Pro Ala glu Asn Pro Ala Pro Ala Glu Ala Asn Val Fhe Leu Gly Thr Gly Ser Ala Asn Val Fhe Leu Gly Fhr Gly Ser Ala Asn Val Ser Asn Val Ser Ser Thr Leu Asn Leu Ser Ser Thr Leu Asn Leu Ser Asn Val Leu Ser Asn Leu Leu Ser Leu Leu<th>Glù Tyr Arg Lys Asn Val Gln Glu Asn Pro Ala Glu Lys 465 Asn Yal Glu Glu Glu Glu 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ggg tct gcc atc cag atg ggg atg gtc ggc atg ctg gar atg ggg atg gag atg gg atg ggg atg at</th>	Glù Tyr Arg Lys Asn Val Gln Glu Asn Pro Ala Glu Lys 465 Asn Yal Glu Glu Glu Glu Glu He Ctt Ctt Gu att Gtt Gu att Asn Val Fee Leu Glu Thr Glu He Ctt Asn July Ser Gu Asn Leu Asn Leu Asn Val Fer Thr Leu Gu Asn Leu Ser Pro 495 aag tca gtg ctc ctg gat ttg ga gac ctt ttg Gu Asn Leu Ser Pro 4495 Asn Leu Leu	agc cag tat cct gas att gtc ttc ctg ggt acg ggg tct gcc atc cag atg ggg atg gtc ggc atg ctg gar atg ggg atg gag atg gg atg ggg atg at

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Le	u Asp	Asr	n Il∈ 100		e Lev	ı Thr	: Arg	Met 105	His 5	s Trp	Sei	r Asr	n Vai	L Gl	y Gly	
Le	u Cys	s Gly 115		: Ile	e Lev	ı Thr	120	Lys	s Glı	ı Thi	r Gl	y Let 12	ı Pro	o Ly	s Cys	

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Ser Gln Tyr Pro Glu Ile Val Phe Leu Gly Thr Gly Ser Ala Ile Pro 470 Met Glu Ile Arg Asn Val Ser Ser Thr Leu Val Asn Leu Ser Pro Asp Lys Ser Val Leu Leu Asp Cys Gly Glu Gly Thr Phe Gly Gln Leu Cys 505 500 Arg His Tyr Gly Gln Gln Ile Asp Arg Val Leu Cys Ser Leu Thr Ala Val Phe Val Ser His Leu His Ala Asp His His Thr Gly Leu Leu Asn 535 Ile Leu Leu Gln Arg Glu His Ala Leu Ala Ser Leu Gly Lys Pro Phe 550 Gln Pro Leu Leu Val Val Ala Pro Thr Gln Leu Arg Ala Trp Leu Gln Gln Tyr His Asn His Cys Gln Glu Ile Leu His His Val Ser Met Ile Pro Ala Lys Cys Leu Gln Lys Gly Ala Glu Val Ser Asn Thr Thr Leu 595 600 Glu Arg Leu Ile Ser Leu Leu Glu Thr Cys Asp Leu Glu Glu Phe Gln Thr Cys Leu Val Arg His Cys Lys His Ala Phe Gly Cys Ala Leu Val His Ser Ser Gly Trp Lys Val Val Tyr Ser Gly Asp Thr Met Pro Cys Glu Ala Leu Val Gln Met Gly Lys Asp Ala Thr Leu Leu Ile His Glu Ala Thr Leu Glu Asp Xaa Leu Glu Glu Glu Ala Val Glu Arg Thr His Ser Thr Thr Ser Gln Ala Ile Asn Val Gly Met Arg Met Asn Ala 695 Glu Phe Ile Met Leu Asn His Phe Ser Gln Arg Tyr Xaa Lys Ile Pro 705 Leu Phe Ser Pro Asp Phe Asn Glu Lys Val Gly Ile Ala Phe Asp His Met Lys Val Xaa Phe Gly Asp Phe Pro Thr Val Pro Lys Leu Ile Pro Pro Leu Lys Ala Leu Phe Ala Gly Asp Ile Glu Glu Met Val Glu Arg Arg Glu Lys Arg Glu Leu Arg Leu Val Arg Ala Ala Leu Leu Thr Gln Gln Ala Asp Ser Pro Glu Asp Arg Glu Pro Gln Gln Lys Arg Ala His

790

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gcg ggt agc cgg Ala Gly Ser Arg 65	gac tcg Asp Ser 70	ggc gcc Gly Ala	gcg c Ala I	ctc tac Leu Tyr 75	gtc tte Val Phe	c tcc e Ser	gag Glu	ttc Phe 80	240
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aag gaa acc ggg Lys Glu Thr Gly 130	ctt cca Leu Pro	aag tgt Lys Cys 135	gta o	ctt tct Leu Ser	gga cc Gly Pr 140	t cca o Pro	caa Gln	ctg Leu	432
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tgc Cys	ctt Leu 610	cag Gln	gaa Glu	ggg ggg	gct Ala	gag Glu 615	atc Ile	tcc Ser	agt Ser	cct Pro	gca Ala 620	gtg Val	gaa Glu	aga Arg	ttg Leu	1872
atc Ile 625	agt Ser	tcg Ser	ctg Leu	ttg Leu	cga Arg 630	aca Thr	tgt Cys	gat Asp	ttg Leu	gaa Glu 635	gag Glu	ttt Phe	cag Gln	acc Thr	tgt Cys 640	1920
ctg Leu	gtg Val	cgg Arg	cac His	tgc Cys 645	aag Lys	cat His	gcg Ala	ttt Phe	ggc Gly 650	Cys	gcg Ala	ctg Leu	gtg Val	cac His 655	Thr	1968
tct Ser	ggc Gly	tgg Trp	aaa Lys 660	Val	gtc Val	tat Tyr	tcc Ser	ggg Gly 665	gac Asp	acc Thr	atg Met	ccc	tgc Cys 670	GLu	gct Ala	2016
ctg Leu	gtc Val	cgg Arg 675	atg Met	Gly	aaa Lys	gat Asp	gcc Ala 680	Thr	ctc Leu	ctg Leu	ata Ile	cat His 685	Glu	gcc Ala	acc Thr	2064
ctg Leu	gaa Glu 690	Asp	ggt Gly	ttg Leu	gaa Glu	gag Glu 695	Glu	gca Ala	gtg Val	gaa Glu	aaq Lys 700	Thr	cac His	ago Ser	aca Thr	2112

acg tcc caa gcc Thr Ser Gln Ala 705	atc agc on Ile Ser V 710	tg ggg atg al Gly Met	cgg atg Arg Met 715	aac gcg gag Asn Ala Glu	ttc att Phe Ile 720	2160
atg ctg aac cad Met Leu Asn His	ttc agc c Phe Ser G 725	ag cgc tat ln Arg Tyr	gcc aag Ala Lys 730	gtc ccc ctc Val Pro Leu	ttc agc Phe Ser 735	2208
ccc aac ttc aac Pro Asn Phe Asr 740	ı Glu Lys V	tg gga gtt al Gly Val 745	gcc ttt Ala Phe	gac cac atg Asp His Met 750	aag gtc Lys Val	2256
tgc ttt gga gad Cys Phe Gly Asp 755	ttt gca a Phe Ala T	ca atg ccc hr Met Pro 760	aag ctg Lys Leu	att ccc cca Ile Pro Pro 765	ctg aaa Leu Lys	2304
gcc ctg ttt gct Ala Leu Phe Ala 770	Gly Asp I	tc gag gag le Glu Glu 75	atg gag Met Glu	gag cgc agg Glu Arg Arg 780	gag aag Glu Lys	2352
cgg gag ctg cgg Arg Glu Leu Arg 785	g cag gtg c g Gln Val A 790	gg gcg gcc rg Ala Ala	ctc ctg Leu Leu 795	tcc agg gag Ser Arg Glu	ctg gca Leu Ala 800	2400
ggc ggc ctg gag Gly Gly Leu Glu	g gat ggg g n Asp Gly G 805	ag cct cag lu Pro Gln	cag aaa Gln Lys 810	cgg gcc cac Arg Ala His	aca gag Thr Glu 815	2448
gag cca cag gcc Glu Pro Gln Ala 820	a Lys Lys V	tc aga gcc al Arg Ala 825	Gln	igatctg ggag	accctg	2498
aattcagaag gct	gtgtgtc tto	tgcccca cg	cacgcacc	cgtatctgcc	ctccttgctg	2558
gtagaagctg aaga	agcacgg tco	cccagga gg	cagctcag	gataggtggt	atggagctgt	2618
gccaaggctt ggg	ctcccac ata	agcacta gt	ctatagat	gcctcttagg	actggtgcct	2678
ggcacagccg cgg	gacagga ggo	tgccaca cg	gaagcaag	cagatgaact	aatttcattt	2738
caaggcagtt ttt	aaagaag gc	tggaaac ag	acggcagc	acctttcctc	taatccagca	2798
aagtgattcc ctg	cacacca gad	gacaagca ga	ıgtaacagg	atcagtgggt	ctaagtgtcc	2858
gagacttaac gaa	aatagta tt	cagctgc aa	taaagatt	gagtttgcaa		2908

<210> 224

<211> 826

<212> PRT

<213> Pan troglodytes

<4.00> 224

Met Trp Ala Leu Cys Ser Leu Leu Arg Ser Ala Ala Gly Arg Thr Met

1 10 15

Ser Gln Gly Arg Thr Ile Ser Gln Ala Pro Ala Arg Arg Glu Arg Pro $20 \\ 25 \\ 30$

Arg Lys Asp Pro Leu Arg His Leu Arg Thr Arg Glu Lys Arg Gly Pro 35 40 45

Ser Gly Cys Ser Gly Gly Pro Asn Thr Val Tyr Leu Gln Val Val Ala Ala Gly Ser Arg Asp Ser Gly Ala Ala Leu Tyr Val Phe Ser Glu Phe Asn Arg Tyr Leu Phe Asn Cys Gly Glu Gly Ile Gln Arg Leu Met Gln Glu His Lys Leu Lys Val Ala Arg Leu Asp Asn Ile Phe Leu Thr Arg Met His Trp Ser Asn Val Gly Gly Leu Ser Gly Met Ile Leu Thr Leu Lys Glu Thr Gly Leu Pro Lys Cys Val Leu Ser Gly Pro Pro Gln Leu Glu Lys Tyr Leu Glu Ala Ile Lys Ile Phe Ser Gly Pro Leu Lys Gly Ile Glu Leu Ala Val Arg Pro His Ser Ala Pro Glu Tyr Glu Asp Glu 170 165 Thr Met Thr Val Tyr Gln Ile Pro Ile His Ser Glu Gln Arg Arg Gly Lys His Gln Pro Trp Gln Ser Pro Glu Arg Pro Leu Ser Arg Leu Ser Pro Glu Arg Ser Ser Asp Ser Glu Ser Asn Glu Asn Glu Pro His Leu Pro His Gly Val Ser Gln Arg Arg Gly Val Arg Asp Ser Ser Leu Val Val Ala Phe Ile Cys Lys Leu His Leu Lys Arg Gly Asn Phe Leu Val Leu Lys Ala Lys Glu Met Gly Leu Pro Val Gly Thr Ala Ala Ile Ala Pro Ile Ile Ala Ala Val Lys Asp Gly Lys Ser Ile Thr His Glu Gly 280 Arg Glu Ile Leu Ala Glu Glu Leu Cys Thr Pro Pro Asp Pro Gly Ala 295 Ala Phe Val Val Val Glu Cys Pro Asp Glu Ser Phe Ile Gln Pro Ile Cys Glu Asn Ala Thr Phe Gln Arg Tyr Gln Gly Lys Ala Asp Ala Pro 325 Val Ala Leu Val Val His Met Ala Pro Glu Ser Val Leu Val Asp Ser 345 Arg Tyr Gln Gln Trp Met Glu Arg Phe Gly Pro Asp Thr Gln His Leu 360 Val Leu Asn Glu Asn Cys Ala Ser Val His Asn Leu Arg Ser His Lys

Ile Gln Thr Gln Leu Asn Leu Ile His Pro Asp Ile Phe Pro Leu Leu 390 395 Thr Ser Phe Pro Cys Lys Lys Glu Gly Pro Thr Leu Ser Val Pro Met Val Gln Gly Glu Cys Leu Leu Lys Tyr Gln Leu Arg Pro Arg Arg Glu Trp Gln Arg Asp Ala Ile Ile Thr Cys Asn Pro Glu Glu Phe Ile Ile 440 Glu Ala Leu Gln Leu Pro Asn Phe Gln Gln Ser Val Gln Glu Tyr Arg Arg Ser Ala Gln Asp Gly Pro Ala Pro Ala Glu Lys Arg Ser Gln Tyr 470 Pro Glu Ile Ile Phe Leu Gly Thr Gly Ser Ala Ile Pro Met Lys Ile 490 Arg Asn Val Ser Ala Thr Leu Val Asn Ile Ser Pro Asp Thr Ser Leu 505 Leu Leu Asp Cys Gly Glu Gly Thr Phe Gly Gln Leu Cys Arg His Tyr 520 Gly Asp Gln Val Asp Arg Val Leu Gly Thr Leu Ala Ala Val Phe Val Ser His Leu His Ala Asp His His Thr Gly Leu Leu Asn Ile Leu Leu Gln Arg Glu Arg Ala Leu Ala Ser Leu Gly Lys Pro Phe His Pro Leu 565 570 Leu Val Val Ala Pro Asn Gln Leu Lys Ala Trp Leu Gln Gln Tyr His 585 Asn Gln Cys Gln Glu Val Leu His His Ile Ser Met Ile Pro Ala Lys Cys Leu Gln Glu Gly Ala Glu Ile Ser Ser Pro Ala Val Glu Arg Leu 615 Ile Ser Ser Leu Leu Arg Thr Cys Asp Leu Glu Glu Phe Gln Thr Cys Leu Val Arg His Cys Lys His Ala Phe Gly Cys Ala Leu Val His Thr 650 Ser Gly Trp Lys Val Val Tyr Ser Gly Asp Thr Met Pro Cys Glu Ala 665 Leu Val Arg Met Gly Lys Asp Ala Thr Leu Leu Ile His Glu Ala Thr Leu Glu Asp Gly Leu Glu Glu Glu Ala Val Glu Lys Thr His Ser Thr 695 Thr Ser Gln Ala Ile Ser Val Gly Met Arg Met Asn Ala Glu Phe Ile

Met	Leu	Asn	His	Phe 725	Ser	Gln	Arg	Tyr	Ala 730	Lys	Val	Pro	Leu	Phe 735	Ser	
Pro	Asn	Phe	Asn 740	Glu	Lys	Val	Gly	Val 745	Ala	Phe	Asp	His	Met 750	Lys	Val	
Cys	Phe	Gly 755	Asp	Phe	Ala	Thr	Met 760	Pro	Lys	Leu	Ile	Pro 765	Pro	Leu	Lys	
Ala	Leu 770	Phe	Ala	Gly	Asp	Ile 775	Glu	Glu	Met	Glu	Glu 780	Arg	Arg	Glu	Lys	
Arg 785	Glu	Leu	Arg	Gln	Val 790	Arg	Ala	Ala	Leu	Leu 795	Ser	Arg	Glu	Leu	Ala 800	
Gly	Gly	Leu	Glu	Asp 805	Gly	Glu	Pro	Gln	Gln 810	Lys	Arg	Ala	His	Thr 815	Glu	
Glu	Pro	Gln	Ala 820	Lys	Lys	Val	Arg	Ala 825	Gln							
<211 <212	0> 22 1> 28 2> DN 3> Go	392	.a go	orill	_a											
	.> CI	os L)	(2478	3)												
atg		25 gcg Ala														48
		gga Gly														96
		gac Asp 35														144
		tgc Cys														192
		agc Ser														240
		tat Tyr														288
		aag Lys														336
atg Met	cac His	tgg Trp 115	tct Ser	aat Asn	gtt Val	ggg Gly	ggc Gly 120	tta Leu	agt Ser	gga Gly	atg Met	att Ile 125	ctt Leu	act Thr	tta Leu	384

Lys G	aa acc lu Thr 30	ggg	ctt Leu	cca Pro	aag Lys 135	tgt Cys	gta Val	ctt Leu	tct Ser	gga Gly 140	cct Pro	cca Pro	cag Gln	ctg Leu	432
gaa aa Glu Ly 145	aa tac ys Tyr	ctc Leu	gaa Glu	gca Ala 150	atc Ile	aaa Lys	ata Ile	ttt Phe	tct Ser 155	ggt Gly	cca Pro	ttg Leu	aaa Lys	gga Gly 160	480
ata ga Ile G	aa ctg lu Leu	gct Ala	gtg Val 165	cgg Arg	ccc Pro	cac His	tct Ser	gcc Ala 170	cca Pro	gaa Glu	tac Tyr	gag Glu	gat Asp 175	gaa Glu	528
acc at Thr Me	tg aca et Thr	gtt Val 180	tac Tyr	cag Gln	atc Ile	ccc Pro	ata Ile 185	cac His	agt Ser	gaa Glu	cag Gln	agg Arg 190	agg Arg	gga Gly	576
agg ca Arg Hi	ac caa is Gln 195	cca Pro	tgg Trp	cag Gln	agt Ser	cca Pro 200	gaa Glu	agg Arg	cct Pro	ctc Leu	agc Ser 205	agg Arg	ctc Leu	agt Ser	624
Pro Gl	ag cga lu Arg 10	tct Ser	tca Ser	gac Asp	tcc Ser 215	gag Glu	tcg Ser	aat Asn	gaa Glu	aat Asn 220	gag Glu	cca Pro	cac His	ctt Leu	672
cca ca Pro Hi 225	at ggt is Gly	gtt Val	agc Ser	cag Gln 230	aga Arg	aga Arg	ggg Gly	gtc Val	agg Arg 235	gac Asp	tct Ser	tcc Ser	ctg Leu	gtc Val 240	720
gta go Val Al	ct ttc la Phe	atc Ile	tgt Cys 245	aag Lys	ctt Leu	cac His	tta Leu	aag Lys 250	aga Arg	gga Gly	aac Asn	ttc Phe	ttg Leu 255	gtg Val	768
ctc aa Leu Ly	aa gca ys Ala	aag Lys 260	gag Glu	atg Met	ggc Gly	ctc Leu	cca Pro 265	gtt Val	Gly	aca Thr	gct Ala	gcc Ala 270	atc Ile	gct Ala	816
ccc at Pro II	tc att le Ile 275	gct Ala	gct Ala	gtc Val	aag Lys	gac Asp 280	ggg Gly	aaa Lys	agc Ser	atc Ile	act Thr 285	cat His	gaa Glu	gga Gly	864
Arg G	ag att lu Ile 90	ttg Leu	gct Ala	gaa Glu	gag Glu 295	ctg Leu	tgt Cys	act Thr	cct Pro	cca Pro 300	gat Asp	cct Pro	ggt Gly	gct Ala	912
gct to Ala Ph 305	tt gtg he Val	gtg Val	gta Val	gaa Glu 310	tgt Cys	cca Pro	gat Asp	gaa Glu	agc Ser 315	ttc Phe	att Ile	caa Gln	ccc Pro	atc Ile 320	960
tgt ga Cys Gi	ag aat lu Asn	gcc Ala	acc Thr 325	ttt Phe	cag Gln	agg Arg	tac Tyr	caa Gln 330	gga Gly	aag Lys	gca Ala	gat Asp	gcc Ala 335	ccc Pro	1008
gtg go Val A	cc ttg la Leu	gtg Val 340	gtt Val	cac His	atg Met	gcc Ala	cca Pro 345	gaa Glu	tct Ser	gtg Val	ctt Leu	gtg Val 350	gac Asp	agc Ser	1056
agg ta Arg T	ac cag yr Gln 355	Gln	tgg Trp	atg Met	gag Glu	agg Arg 360	ttt Phe	ggg Gly	cct Pro	gac Asp	acc Thr 365	cag Gln	cac His	ttg Leu	1104
Val L	tg aat eu Asn 70	gag Glu	aac Asn	tgt Cys	gcc Ala 375	tca Ser	gtt Val	cac His	aac Asn	ctt Leu 380	cgc Arg	agc Ser	cac His	aag Lys	1152

			-		aac Asn 390				_	_				-		1200
					aag Lys											1248
					ctc Leu											1296
					att Ile											1344
		_	_		ccc Pro			-	_	_		-				1392
					gtc Val 470											1440
					ctt Leu											1488
cga Arg	aat Asn	gtc Val	agt Ser 500	gcc Ala	aca Thr	ctt Leu	gtc Val	aac Asn 505	ata Ile	agc Ser	ccc Pro	gac Asp	acg Thr 510	tct Ser	ctg Leu	1536
cta Leu	ctg Leu	gac Asp 515	tgt Cys	ggt Gly	gag Glu	ggc Gly	acg Thr 520	ttt Phe	ggg Gly	cag Gln	ctg Leu	tgc Cys 525	cgt Arg	cat His	tac Tyr	1584
gga Gly	gac Asp 530	cag Gln	gtg Val	gac Asp	agg Arg	gtc Val 535	ctg Leu	ggc Gly	acc Thr	ctg Leu	gct Ala 540	gct Ala	gtg Val	ttt Phe	gtg Val	1632
tcc Ser 545	cac His	ctg Leu	cac His	gca Ala	gat Asp 550	cac His	cac His	acg Thr	ggc Gly	ttg Leu 555	cta Leu	aat Asn	atc Ile	țtg Leu	ctg Leu 560	1680
					ttg Leu											1728
					agc Ser											1776
aac Asn	cag Gln	tgc Cys 595	cag Gln	gag Glu	gtc Val	ctg Leu	cac His 600	cac His	atc Ile	agt Ser	atg Met	att Ile 605	cct Pro	gcc Ala	aaa Lys	1824
tgc Cys	ctt Leu 610	cag Gln	gaa Glu	ggg Gly	gct Ala	gag Glu 615	atc Ile	tcc Ser	agt Ser	cct Pro	gca Ala 620	gtg Val	gaa Glu	aga Arg	ttg Leu	1872
atc Ile 625	agt Ser	tcg Ser	ctg Leu	ttg Leu	cga Arg 630	aca Thr	tgt Cys	gat Asp	ttg Leu	gaa Glu 635	gag Glu	ttt Phe	cag Gln	acc Thr	tgt Cys 640	1920

														cac His 655		1968
														gag Glu		2016
														gcc Ala		2064
														agc Ser		2112
														ttc Phe		2160
atg Met	ctg Leu	aac Asn	cac His	ttc Phe 725	agc Ser	cag Gln	cgc Arg	tat Tyr	gcc Ala 730	aag Lys	gtc Val	ccc Pro	ctc Leu	ttc Phe 735	agc Ser	2208
ccc Pro	aac Asn	ttc Phe	aac Asn 740	gag Glu	aaa Lys	gtg Val	gga Gly	gtt Val 745	gcc Ala	ttt Phe	gac Asp	cac	atg Met 750	aag Lys	gtc Val	2256
														ctg Leu		2304
														gag Glu		2352
cgg Arg 785	gag Glu	ctg Leu	cgg Arg	cag Gln	gtg Val 790	cgg Arg	gcg Ala	gcc Ala	ctc Leu	ctg Leu 795	tcc Ser	Gly ggg	gag Glu	ctg Leu	gca Ala 800	2400
														aca Thr 815		2448
		cag Gln								tgaa	agato	ctg (ggag	accct	tg	2498
aatt	caga	ag ç	gctgt	gtgt	c tt	ctg	ccca	a cgo	cacgo	cacc	cgt	atct	gcc	ctcc [.]	ttgctg	2558
gtag	gaago	ctg a	aagaq	gcac	gg to	ccc	cagga	a ggo	cagct	cag	gat	aggt	ggt	atgg	agctgt	2618
gccg	gaggo	ctt a	aggct	ccca	ac at	caago	cacta	a gto	ctata	aggt	gcc	tggc	aca	gccg	cgggac	2678
agga	aggct	gc d	cacac	cggaa	ag ca	aagca	agato	g aa	ctaat	tttc	att	tcaa	ggc	agtt	tttaaa	2738
,gaaq	gtctt	gg a	aaaca	agaco	gg ca	agca	cctt	t cct	tctaa	atcc	agc	aaag	tga	ttcc	ctgcac	2798
acca	agaga	aca a	agcaç	gagta	aa ca	agga	tcact	t gg	gtcta	aagt	gtc	cgag	act	taac	gaaaat	2858
agta	attto	cag o	ctgca	aataa	aa ga	attga	agtti	t gc	aa							2892

<210> 226 <211> 826 <212> PRT

<213> Gorilla gorilla

 $<\!400\!>$ 226 Met Trp Ala Leu Cys Ser Leu Leu Arg Ser Ala Ala Gly Arg Thr Met 1 5 10 15

Ser Gln Gly Arg Thr Ile Ser Gln Ala Pro Ala Arg Arg Glu Arg Pro 20 25 30

Arg Lys Asp Pro Leu Arg His Leu Arg Thr Arg Glu Lys Arg Gly Pro $35 \hspace{1cm} 40 \hspace{1cm} 45$

Ser Gly Cys Ser Gly Gly Pro Asn Thr Val Tyr Leu Gln Val Val Ala 50 55 60

Ala Gly Ser Arg Asp Ser Gly Ala Ala Leu Tyr Val Phe Ser Glu Phe 65 70 75 80

Asn Arg Tyr Leu Phe Asn Cys Gly Glu Gly Val Gln Arg Leu Met Gln 85 90 95

Glu His Lys Leu Lys Val Val Arg Leu Asp Asn Ile Phe Leu Thr Arg 100 105 110

Met His Trp Ser Asn Val Gly Gly Leu Ser Gly Met Ile Leu Thr Leu 115 120 125

Lys Glu Thr Gly Leu Pro Lys Cys Val Leu Ser Gly Pro Pro Gln Leu 130 135 140

Glu Lys Tyr Leu Glu Ala Ile Lys Ile Phe Ser Gly Pro Leu Lys Gly 145 150 155 160

Ile Glu Leu Ala Val Arg Pro His Ser Ala Pro Glu Tyr Glu Asp Glu 165 170 175

Thr Met Thr Val Tyr Gln Ile Pro Ile His Ser Glu Gln Arg Arg Gly 180 185 190

Arg His Gln Pro Trp Gln Ser Pro Glu Arg Pro Leu Ser Arg Leu Ser 195 200 205

Pro Glu Arg Ser Ser Asp Ser Glu Ser Asn Glu Asn Glu Pro His Leu 210 215 220

Pro His Gly Val Ser Gln Arg Arg Gly Val Arg Asp Ser Ser Leu Val 225 230 235 240

Val Ala Phe Ile Cys Lys Leu His Leu Lys Arg Gly Asn Phe Leu Val 245 250 255

Leu Lys Ala Lys Glu Met Gly Leu Pro Val Gly Thr Ala Ala Ile Ala 260 265 270

Pro Ile Ile Ala Ala Val Lys Asp Gly Lys Ser Ile Thr His Glu Gly 275 280 285

Arg Glu Ile Leu Ala Glu Glu Leu Cys Thr Pro Pro Asp Pro Gly Ala 290 295 300

Ala Phe Val Val Val Glu Cys Pro Asp Glu Ser Phe Ile Gln Pro Ile 315 Cys Glu Asn Ala Thr Phe Gln Arg Tyr Gln Gly Lys Ala Asp Ala Pro 330 Val Ala Leu Val Val His Met Ala Pro Glu Ser Val Leu Val Asp Ser Arg Tyr Gln Gln Trp Met Glu Arg Phe Gly Pro Asp Thr Gln His Leu Val Leu Asn Glu Asn Cys Ala Ser Val His Asn Leu Arg Ser His Lys 375 Ile Gln Thr Gln Leu Asn Leu Ile His Pro Asp Ile Phe Pro Leu Leu Thr Ser Phe Pro Cys Lys Lys Glu Gly Pro Thr Leu Ser Val Pro Met Val Gln Gly Glu Cys Leu Leu Lys Tyr Gln Leu Arg Pro Arg Arg Glu Trp Gln Arg Asp Ala Ile Ile Thr Cys Asn Pro Glu Glu Phe Ile Val Glu Ala Leu Gln Leu Pro Asn Phe Gln Gln Ser Val Gln Glu Tyr Arg Arg Ser Val Gln Asp Val Pro Ala Pro Ala Glu Lys Arg Ser Gln Tyr 470 Pro Glu Ile Ile Phe Leu Gly Thr Gly Ser Ala Ile Pro Met Lys Ile Arg Asn Val Ser Ala Thr Leu Val Asn Ile Ser Pro Asp Thr Ser Leu 505 Leu Leu Asp Cys Gly Glu Gly Thr Phe Gly Gln Leu Cys Arg His Tyr Gly Asp Gln Val Asp Arg Val Leu Gly Thr Leu Ala Ala Val Phe Val 535 Ser His Leu His Ala Asp His His Thr Gly Leu Leu Asn Ile Leu Leu Gln Arg Glu Gln Ala Leu Ala Ser Leu Gly Lys Pro Leu His Pro Leu Leu Val Val Ala Pro Ser Gln Leu Lys Ala Trp Leu Gln Gln Tyr His Asn Gln Cys Gln Glu Val Leu His His Ile Ser Met Ile Pro Ala Lys 595 600 Cys Leu Gln Glu Gly Ala Glu Ile Ser Ser Pro Ala Val Glu Arg Leu 615 Ile Ser Ser Leu Leu Arg Thr Cys Asp Leu Glu Glu Phe Gln Thr Cys 630 635

 Leu
 Val
 Arg
 His
 Cys
 His
 Ala
 Phe
 Gly
 Cys
 Ala
 Leu
 Val
 His
 Thr

 Ser
 Gly
 Trp
 Lys
 Val
 Tyr
 Ser
 Gly
 Asp
 Thr
 Met
 Pro
 Cys
 Glu
 Ala

 Leu
 Val
 Asp
 Gly
 Lys
 Asp
 Ala
 Thr
 Leu
 Leu
 His
 Glu
 Ala
 Thr

 Leu
 Gly
 Asp
 Gly
 Lys
 Gly
 Asp
 Ala
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 Fro
 Asp
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<210> 227

<211> 844

<212> PRT

<213> Caenorhabditis elegans

<400> 227

Met Lys Met Leu Phe Phe Gly Ile Lys Val Ser Arg His Leu Ile Ser 1 5 10 15

Ser Thr Ser Cys Leu Phe Lys Asp Asn Asn Glu Glu Leu Leu Glu Ser 20 25 30

Ile Lys Glu Arg Ile Ala Arg Asn Arg Arg Ile Leu Gln Lys His Ser 35 40 45

Ser Ser His Leu Lys Ala Arg Glu Val Asn Ala Ser Ile Ser Asn Leu 50 55 60

Arg Gln Ser Met Ala Ala Val Gln Lys Lys Gln Lys Ala Ala His Glu
65 70 75 80

Pro Pro Ala Asn Ser Ile Val Asn Ile Pro Ser Gln Val Ser Ile Glu 85 90 95

Val Leu Gly Asn Gly Thr Gly Leu Leu Arg Ala Cys Phe Ile Leu Arg 100 105 110 Thr Pro Leu Lys Thr Tyr Met Phe Asn Cys Pro Glu Asn Ala Cys Arg 120 Phe Leu Trp Gln Leu Arg Ile Arg Ser Ser Ser Val Val Asp Leu Phe 135 Ile Thr Ser Ala Asn Trp Asp Asn Ile Ala Gly Ile Ser Ser Ile Leu Leu Ser Lys Glu Ser Asn Ala Leu Ser Thr Arg Leu His Gly Ala Met 170 Asn Ile Lys His Phe Leu Glu Cys Ile Arg Pro Phe Gln Asp Ser Asp Tyr Gly Ser Cys Lys Tyr Pro Ser Gln Val Glu Glu Arg Pro Tyr Thr Met Glu Asn Tyr Glu Asp Ala Gly Leu Lys Val Thr Tyr Ile Pro Leu 215 Ser Pro Pro Leu Asn Ile Gly Ser Asn Asn Glu Lys Ser Lys Asn Val Lys Val Asn Asn Val Asp Ile Ala Phe Leu Ile Glu Met Lys Glu Ala Ala Arg Arg Ile Asp Thr Met Lys Leu Met Glu Leu Lys Val Pro Lys Gly Pro Leu Ile Gly Lys Leu Lys Ser Gly Glu Ala Val Thr Leu Pro Asp Gly Arg Thr Ile Gln Pro Asp Gln Val Phe Ser Ser Asp Lys Val Glu Gly Asp Lys Pro Leu Leu Leu Val Thr Glu Cys Thr Thr Glu Asp 315 310 His Val Lys Ala Leu Ile Asp Ser Ser Ser Leu Gln Pro Phe Leu Asn Gly Glu Lys Gln Leu Asp Tyr Met Val His Ile Ser Asp Asp Ala Val 345 Ile Asn Thr Pro Thr Tyr Arg His Leu Met Glu Lys Leu Asn Asn Pro 360 Ser Ile Thr His Leu Leu Ile Asn Gly Gly Asn Pro Val Ile Pro Ala Val Glu Ser Val Tyr Lys His Thr Arg Leu Leu Arg Ser Ile Ala Pro Ser Leu Phe Pro Ala Leu His Pro Ile Asp Trp Ser Gly Ile Ile Thr 410 405 Gln Asn Glu Glu Leu Ser Gln Arg Gln Asp Gln Phe Ile Arg Val Ala 425 Pro Met Gln Arg Tyr Trp Met Arg Arg Gly Ala Ser Phe Asn Glu Glu Pro Ile Val Asn Asn Leu Leu Ala Ala Glu Pro Glu Leu Ser Asp Lys Ala Lys Glu Leu Ile Lys Glu Tyr Gln Lys Leu Glu Lys Glu Asn Lys Met Asp Cys Glu Phe Pro Lys Leu Thr Phe Phe Gly Thr Ser Ser Ala 490 Val Pro Ser Lys Tyr Arg Asn Val Thr Gly Tyr Leu Val Glu Ala Ser Glu Asn Ser Ala Ile Leu Ile Asp Val Gly Glu Gly Thr Tyr Gly Gln Met Arg Ala Val Phe Gly Glu Asp Gly Cys Lys Gln Leu Leu Val Asn 535 Leu Asn Cys Val Leu Ile Thr His Ala His Gln Asp His Met Asn Gly Leu Tyr Thr Ile Ile Ala Arg Arg Lys Glu Ala Phe Glu Ser Leu Gly 570 Ala Pro Tyr Arg Pro Leu Val Leu Val Cys Asn Arg Asn Val Leu Lys Pro Met Lys Thr Tyr Ser Ile Cys Phe Glu Asn Ile Glu His Leu Leu 600 Glu Ile Val Asp Ile Ser Arg Tyr Pro Leu Thr Pro Pro Gly Ser Pro Gly Gly Pro Pro Gly Lys Arg Pro Arg Leu Pro Ser Pro His Leu Pro Pro Ser Arg Asp Val Leu Gln Asp Met Ser Ser Phe Asp Lys Lys Ala Trp Lys Leu Asp Glu Leu Lys Ala Val Gln Val His His Thr Arg 665 Met Ala Asn Gly Phe Val Met Arg Val Ala Gly Lys Arg Ile Val Phe Ser Gly Asp Thr Lys Pro Cys Asp Leu Leu Val Glu Glu Gly Lys Asp 695 Ala Asp Val Leu Val His Glu Ser Thr Phe Glu Asp Gly His Glu Val 715 710 Asp Met Thr Pro Lys Pro Pro Lys Lys Leu Ala Lys Ile Ser Ser Leu Ala Asp Ala Met Arg Lys Arg His Ser Thr Met Gly Gln Ala Val Asp Val Gly Lys Arg Met Asn Ala Lys His Ile Ile Leu Thr His Phe Ser 760 Ala Arg Tyr Pro Lys Val Pro Val Leu Pro Glu Tyr Leu Asp Lys Glu Asn Ile Gly Val Ala Met Asp Met Leu Arg Val Arg Phe Asp His Leu 785 790 795 800

Pro Leu Val Ser Lys Leu Pro Ile Phe Arg Glu Val Phe Val Ala 805 810 815

Glu Leu Phe Glu Leu Thr Ile Lys Lys Glu Gln Arg Val Leu Lys Asp 820 825 830

Lys Glu Leu Ser Glu Lys Arg Gly Gln Leu Lys Ala 835 840

<210> 228

<211> 837

<212> PRT

<213> Arabidopsis thaliana

<400> 228

Met Glu Asn Asn Glu Ala Thr Asn Gly Ser Lys Ser Ser Ser Asn Ser 1 15

Phe Val Phe Asn Lys Arg Arg Ala Glu Gly Phe Asp Ile Thr Asp Lys 20 25 30

Lys Lys Arg Asn Leu Glu Arg Lys Ser Gln Lys Leu Asn Pro Thr Asn 35 40 45

Thr Ile Ala Tyr Ala Gln Ile Leu Gly Thr Gly Met Asp Thr Gln Asp 50 55 60

Thr Ser Ser Ser Val Leu Leu Phe Phe Asp Lys Gln Arg Phe Ile Phe 65 70 75 80

Asn Ala Gly Glu Gly Leu Gln Arg Phe Cys Thr Glu His Lys Ile Lys 85 90 95

Leu Ser Lys Ile Asp His Val Phe Leu Ser Arg Val Cys Ser Glu Thr 100 105 110

Ala Gly Gly Leu Pro Gly Leu Leu Leu Thr Leu Ala Gly Ile Gly Glu 115 120 125

Glu Gly Leu Ser Val Asn Val Trp Gly Pro Ser Asp Leu Asn Tyr Leu 130 135

Val Asp Ala Met Lys Ser Phe Ile Pro Arg Ala Ala Met Val His Thr 145 150 155 160

Arg Ser Phe Gly Pro Ser Ser Thr Pro Asp Pro Ile Val Leu Val Asn 165 170 175

Asp Glu Val Val Lys Ile Ser Ala Ile Ile Leu Lys Pro Cys His Ser 180 185 190

Glu Glu Asp Ser Gly Asn Lys Ser Gly Asp Leu Ser Val Val Tyr Val 195 200 205

Cys Glu Leu Pro Glu Ile Leu Gly Lys Phe Asp Leu Glu Lys Ala Lys 210 215 220

Lys Val Phe Gly Val Lys Pro Gly Pro Lys Tyr Ser Arg Leu Gln Ser 225 230 235

Gly Glu Ser Val Lys Ser Asp Glu Arg Asp Ile Thr Val His Pro Ser Asp Val Met Gly Pro Ser Leu Pro Gly Pro Ile Val Leu Leu Val Asp Cys Pro Thr Glu Ser His Ala Ala Glu Leu Phe Ser Leu Lys Ser Leu 280 Glu Ser Tyr Tyr Ser Ser Pro Asp Glu Gln Thr Ile Gly Ala Lys Phe Val Asn Cys Ile Ile His Leu Ser Pro Ser Ser Val Thr Ser Ser Pro Thr Tyr Gln Ser Trp Met Lys Lys Phe His Leu Thr Gln His Ile Leu 330 Ala Gly His Gln Arg Phe Leu Pro Leu Leu Ile Ile Val Ser His Gln Lys Thr Val Arg Lys Asn Met Ala Phe Pro Ile Leu Lys Ala Ser Ser 360 Arg Ile Ala Ala Arg Leu Asn Tyr Leu Cys Pro Gln Phe Pro Ala Pro Gly Phe Trp Pro Ser Gln Leu Thr Asp Asn Ser Ile Ile Asp Pro 390 Thr Pro Ser Asn Lys Phe Asn Leu Arg Pro Val Ala Ile Arg Gly Ile Asp Arg Ser Cys Ile Pro Ala Pro Leu Thr Ser Ser Glu Val Val Asp Glu Leu Leu Ser Glu Ile Pro Glu Ile Lys Asp Lys Ser Glu Glu Ile Lys Gln Phe Trp Asn Lys Gln His Asn Lys Thr Ile Ile Glu Lys Leu Trp Leu Ser Glu Cys Asn Thr Val Leu Pro Asn Cys Leu Glu Lys Ile 475 Arg Arg Asp Asp Met Glu Ile Val Ile Leu Gly Thr Gly Ser Ser Gln Pro Ser Lys Tyr Arg Asn Val Ser Ala Ile Phe Ile Asp Leu Phe Ser 505 Arg Gly Ser Leu Leu Leu Asp Cys Gly Glu Gly Thr Leu Gly Gln Leu Lys Arg Arg Tyr Gly Leu Asp Gly Ala Asp Glu Ala Val Arg Lys Leu Arg Cys Ile Trp Ile Ser His Ile His Ala Asp His His Thr Gly Leu 555 550 Ala Arg Ile Leu Ala Leu Arg Ser Lys Leu Leu Lys Gly Val Thr His

Glu Pro Val Ile Val Val Gly Pro Arg Pro Leu Lys Arg Phe Leu Asp 585 Ala Tyr Gln Arg Leu Glu Asp Leu Asp Met Glu Phe Leu Asp Cys Arg Ser Thr Thr Ala Thr Ser Trp Ala Ser Leu Glu Ser Gly Gly Glu Ala Glu Gly Ser Leu Phe Thr Gln Gly Ser Pro Met Gln Ser Val Phe Lys Arg Ser Asp Ile Ser Met Asp Asn Ser Ser Val Leu Leu Cys Leu Lys 645 655 Asn Leu Lys Lys Val Leu Ser Glu Ile Gly Leu Asn Asp Leu Ile Ser 665 Phe Pro Val Val His Cys Pro Gln Ala Tyr Gly Val Val Ile Lys Ala Ala Glu Arg Val Asn Ser Val Gly Glu Gln Ile Leu Gly Trp Lys Met 695 Val Tyr Ser Gly Asp Ser Arg Pro Cys Pro Glu Thr Val Glu Ala Ser Arg Asp Ala Thr Ile Leu Ile His Glu Ala Thr Phe Glu Asp Ala Leu Ile Glu Glu Ala Leu Ala Lys Asn His Ser Thr Thr Lys Glu Ala Ile Asp Val Gly Ser Ala Ala Asn Val Tyr Arg Ile Val Leu Thr His Phe 760 Ser Gln Arg Tyr Pro Lys Ile Pro Val Ile Asp Glu Ser His Met His Asn Thr Cys Ile Ala Phe Asp Leu Met Ser Ile Asn Met Ala Asp Leu His Val Leu Pro Lys Val Leu Pro Tyr Phe Lys Thr Leu Phe Arg Asp 810 Glu Met Val Glu Asp Glu Asp Ala Asp Asp Val Ala Met Asp Asp Leu Lys Glu Glu Ala Leu 835

<210> 229

<211> 838

<212> PRT

<213> Saccharomyces cerevisiae

<400> 229

Met Phe Thr Phe Ile Pro Ile Thr His Pro Thr Ser Asp Thr Lys His

1 10 15

Pro Leu Leu Val Gln Ser Ala His Gly Glu Lys Tyr Phe Phe Gly

Lys Ile Gly Glu Gly Ser Gln Arg Ser Leu Thr Glu Asn Lys Ile Arg Ile Ser Lys Leu Lys Asp Ile Phe Leu Thr Gly Glu Leu Asn Trp Ser Asp Ile Gly Gly Leu Pro Gly Met Ile Leu Thr Ile Ala Asp Gln Gly Lys Ser Asn Leu Val Leu His Tyr Gly Asn Asp Ile Leu Asn Tyr Ile Val Ser Thr Trp Arg Tyr Phe Val Phe Arg Phe Gly Ile Asp Leu Asn Asp His Ile Met Lys Asp Lys Glu Val Tyr Lys Asp Lys Ile Ile Ala Val Lys Ser Phe Asn Val Leu Lys Asn Gly Gly Glu Asp Arg Leu Gly Val Phe Asp Ser Phe Gln Lys Gly Val Leu Arg Ser Ile Val Ala Lys 150 Met Phe Pro Lys His Ala Pro Thr Asp Arg Tyr Asp Pro Ser Ser Asp Pro His Leu Asn Val Glu Leu Pro Asp Leu Asp Ala Lys Val Glu Val Ser Thr Asn Tyr Glu Ile Ser Phe Ser Pro Val Arg Gly Lys Phe Lys 200 Val Glu Glu Ala Ile Lys Leu Gly Val Pro Lys Gly Pro Leu Phe Ala Lys Leu Thr Lys Gly Gln Thr Ile Thr Leu Asp Asn Gly Ile Val Val 235 Thr Pro Glu Gln Val Leu Glu Asn Glu Arg His Phe Ala Lys Val Leu Ile Leu Asp Ile Pro Asp Asp Leu Tyr Leu Asn Ala Phe Val Glu Lys 265 Phe Lys Asp Tyr Asp Cys Ala Glu Leu Gly Met Val Tyr Tyr Phe Leu Gly Asp Glu Val Thr Ile Asn Asp Asn Leu Phe Ala Phe Ile Asp Ile 295 Phe Glu Lys Asn Asn Tyr Gly Lys Val Asn His Met Ile Ser His Asn Lys Ile Ser Pro Asn Thr Ile Ser Phe Phe Gly Ser Ala Leu Thr Thr 335 Leu Lys Leu Lys Ala Leu Gln Val Asn Asn Tyr Asn Leu Pro Lys Thr 340 345 Asp Arg Val Phe Ser Lys Asp Phe Tyr Asp Arg Phe Asp Thr Pro Leu

Ser Arg Gly Thr Ser Met Cys Lys Ser Gln Glu Glu Pro Leu Asn Thr Ile Ile Glu Lys Asp Asn Ile His Ile Phe Ser Gln Asn Lys Thr Val 395 390 Thr Phe Glu Pro Phe Arg Met Asn Glu Glu Pro Met Lys Cys Asn Ile Asn Gly Glu Val Ala Asp Phe Ser Trp Gln Glu Ile Phe Glu Glu His 425 Val Lys Pro Leu Glu Phe Pro Leu Ala Asp Val Asp Thr Val Ile Asn 440 Asn Gln Leu His Val Asp Asn Phe Asn Asn Ser Ala Glu Lys Lys 455 His Val Glu Ile Ile Thr Leu Gly Thr Gly Ser Ala Leu Pro Ser Lys Tyr Arg Asn Val Val Ser Thr Leu Val Lys Val Pro Phe Thr Asp Ala Asp Gly Asn Thr Ile Asn Arg Asn Ile Met Leu Asp Ala Gly Glu Asn Thr Leu Gly Thr Ile His Arg Met Phe Ser Gln Leu Ala Val Lys Ser Ile Phe Gln Asp Leu Lys Met Ile Tyr Leu Ser His Leu His Ala Asp 535 His His Leu Gly Ile Ile Ser Val Leu Asn Glu Trp Tyr Lys Tyr Asn Lys Asp Asp Glu Thr Ser Tyr Ile Tyr Val Val Thr Pro Trp Gln Tyr His Lys Phe Val Asn Glu Trp Leu Val Leu Glu Asn Lys Glu Ile Leu Lys Arg Ile Lys Tyr Ile Ser Cys Glu His Phe Ile Asn Asp Ser Phe Val Arg Met Gln Thr Gln Ser Val Pro Leu Ala Glu Phe Asn Glu Ile Leu Lys Glu Asn Ser Asn Gln Glu Ser Asn Arg Lys Leu Glu Leu Asp Arg Asp Ser Ser Tyr Arg Asp Val Asp Leu Ile Arg Gln Met Tyr Glu Asp Leu Ser Ile Glu Tyr Phe Gln Thr Cys Arg Ala Ile His Cys Asp Trp Ala Tyr Ser Asn Ser Ile Thr Phe Arg Met Asp Glu Asn Asn Glu 685 680 His Asn Thr Phe Lys Val Ser Tyr Ser Gly Asp Thr Arg Pro Asn Ile

Glu Lys Phe Ser Leu Glu Ile Gly Tyr Asn Ser Asp Leu Leu Ile His 705 710 715 720

Glu Ala Thr Leu Glu Asn Gln Leu Leu Glu Asp Ala Val Lys Lys 725 730 730 735

His Cys Thr Ile Asn Glu Ala Ile Gly Val Ser Asn Lys Met Asn Ala 740 745 750 ·

Arg Lys Leu Ile Leu Thr His Phe Ser Gln Arg Tyr Pro Lys Leu Pro
755 760 765

Gln Leu Asp Asn Asn Ile Asp Val Met Ala Arg Glu Phe Cys Phe Ala 770 780

Phe Asp Ser Met Ile Val Asp Tyr Glu Lys Ile Gly Glu Gln Gln Arg 785 790 795 800

Ile Phe Pro Leu Leu Asn Lys Ala Phe Val Glu Glu Lys Glu Glu Glu 805 810 815

Glu Asp Val Asp Asp Val Glu Ser Val Gln Asp Leu Glu Val Lys Leu 820 825 830

Lys Lys His Lys Lys Asn 835

<210> 230

<211> 311

<212> PRT

<213> Escherichia coli

<400> 230

Met Lys Arg Asp Glu Leu Met Glu Leu Ile Phe Leu Gly Thr Ser Ala 1 5 10 15

Gly Val Pro Thr Arg Thr Arg Asn Val Thr Ala Ile Leu Leu Asn Leu 20 25 30

Gln His Pro Thr Gln Ser Gly Leu Trp Leu Phe Asp Cys Gly Glu Gly 35 40 45

Thr Gln His Gln Leu Leu His Thr Ala Phe Asn Pro Gly Lys Leu Asp 50 55 60

Lys Ile Phe Ile Ser His Leu His Gly Asp His Leu Phe Gly Leu Pro 65 70 75 80

Gly Leu Cys Ser Arg Ser Met Ser Gly Ile Ile Gln Pro Leu Thr 85 90 95

Ile Tyr Gly Pro Gln Gly Ile Arg Glu Phe Val Glu Thr Ala Leu Arg 100 105 110

Ile Ser Gly Ser Trp Thr Asp Tyr Pro Leu Glu Ile Val Glu Ile Gly 115 120 125

Ala Gly Glu Ile Leu Asp Asp Gly Leu Arg Lys Val Thr Ala Tyr Pro 130 140

Leu Glu His Pro Leu Glu Cys Tyr Gly Tyr Arg Ile Glu Glu His Asp 145 150 155 160

Pro Gly Pro Leu Phe Gln Glu Leu Lys Ala Gly Lys Thr Ile Thr Leu 180 185 190

Glu Asp Gly Arg Gln Ile Asn Gly Ala Asp Tyr Leu Ala Ala Pro Val 195 200 205

Pro Gly Lys Ala Leu Ala Ile Phe Gly Asp Thr Gly Pro Cys Asp Ala 210 215 220

Ala Leu Asp Leu Ala Lys Gly Val Asp Val Met Val His Glu Ala Thr 225 230 235 240

Leu Asp Ile Thr Met Glu Ala Lys Ala Asn Ser Arg Gly His Ser Ser 245 250 255

Thr Arg Gln Ala Ala Thr Leu Ala Arg Glu Ala Gly Val Gly Lys Leu 260 265 270

Ile Ile Thr His Val Ser Ser Arg Tyr Asp Asp Lys Gly Cys Gln His 275 280 285

Leu Leu Arg Glu Cys Arg Ser Ile Phe Pro Ala Thr Glu Leu Ala Asn 290 295 300

Asp Phe Thr Val Phe Asn Val 305 310

<210> 231

<211> 326

<212> PRT

<213> Synechocystis sp.

<400> 231

Met Glu Ile Thr Phe Leu Gly Thr Ser Ser Gly Val Pro Thr Arg Asn
1 5 10 15

Trp Leu Phe Asp Cys Gly Glu Gly Thr Gln His Gln Phe Leu Arg Ser

Glu Val Lys Ile Ser Gln Leu Thr Arg Ile Phe Ile Thr His Leu His 50 55 60

Gly Asp His Ile Phe Gly Leu Met Gly Leu Leu Ala Ser Ser Gly Leu 65 70 75 80

Ala Gly Ser Gly Gln Gly Ile Glu Ile Tyr Gly Pro Glu Gly Leu Gly 85 90 95

Asp Tyr Leu Glu Ala Cys Cys Arg Phe Ser Ser Thr His Leu Gly Lys 100 105 110

Arg Leu Lys Val His Thr Val Arg Glu Asn Gly Leu Ile Tyr Glu Asp 115 120 125

Lys Asp Phe Gln Val His Cys Gly Leu Leu Lys His Arg Ile Pro Ala 130 135 140

Tyr Gly Tyr Arg Val Glu Glu Lys Gln Arg Pro Gly Arg Phe Asn Val Glu Gln Ala Glu Ala Leu Gly Ile Pro Phe Gly Pro Ile Tyr Gly Gln Leu Lys Gln Gly Lys Thr Val Thr Leu Glu Asp Gly Arg Arg Ile Arg 185 Gly Gln Asp Leu Cys Glu Pro Pro Glu Pro Gly Arg Lys Phe Val Tyr 200 Cys Thr Asp Thr Val Phe Cys Glu Glu Ala Ile Ala Leu Ala Gln Glu Ala Asp Leu Leu Val His Glu Ala Thr Phe Ala His Gln Asp Ala Gln Leu Ala Phe Asp Arg Leu His Ser Thr Ser Thr Met Ala Ala Gln Val 250 Ala Leu Leu Ala Asn Val Lys Gln Leu Ile Met Thr His Phe Ser Pro Arg Tyr Ala Pro Gly Asn Pro Leu Gln Leu Glu Asn Leu Leu Ala Glu Ala Gln Ala Ile Phe Pro Asn Thr Arg Leu Ala Arg Asp Phe Leu Thr 295 300 Val Glu Ile Pro Arg Arg Thr Ala Asp Pro Ala Ile Ala Met Ser Thr 310 Pro Gln Ala Ser Pro Ala

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<210> 232
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<400> 232

Met Met Glu Val Thr Phe Leu Gly Thr Ser Ser Ala Val Pro Ser Lys
1 5 10 15

Asn Arg Asn His Thr Ser Ile Ala Leu Arg Ile Pro Gly Glu Ile Phe 20 25 30

Leu Phe Asp Cys Gly Glu Gly Thr Gln Arg Gln Met Ala Leu Ala Gly 35 40 45

Ile Ser Pro Met Lys Val Thr Arg Ile Phe Ile Thr His Leu His Gly
50 55 60

Asp His Ile Leu Gly Ile Pro Gly Met Ile Gln Ser Met Gly Phe Arg
65 70 75 80

Gly Arg Glu Glu Pro Leu Asp Ile Tyr Gly Pro Pro Gly Ile His Glu 85 90 95

Leu His Glu Cys Ile Met Lys Met Gly Tyr Phe Thr Leu Asp Phe Asp 100 105 110

<211> 307

<212> PRT

<213> Methanobacterium thermoautotrophicum

lle Asn Val His Glu Val Arg Gly Gly Thr Val Val Glu Glu Asp Asp 115 120 125

Tyr Arg Val Thr Ser Ala Pro Ala Ser His Ser Val Phe Asn Leu Ala 130 135 140

Tyr Cys Phe Glu Glu Lys Lys Arg Pro Arg Phe Leu Arg Glu Lys Ala 145 150 155 160

Ile Ala Leu Gly Leu Lys Pro Gly Pro Ala Phe Gly Lys Leu His Arg 165 170 175

Gly Ile Pro Val Arg Val Gly Asp Arg Ile Ile Met Pro Glu Glu Val 180 185 190

Leu Gly Ser Pro Arg Lys Gly Val Lys Val Cys Tyr Ser Gly Asp Thr 195 200 205

Arg Pro Cys Glu Ser Val Ile Lys Leu Ala Glu Gly Ala Glu Leu Leu 210 215 220

Ile His Glu Ser Thr Leu Glu Ala Gly Ser Glu Asp Lys Ala Ala Glu 225 230 235 240

Ser Gly His Ser Thr Ala Arg Glu Ala Ala Glu Val Ala Arg Ser Ala 245 250 255

Gly Val Lys Arg Leu Ile Leu Thr His Leu Ser Thr Arg Tyr Lys Arg 260 265 270

Thr Glu Val Ile Leu Glu Ala Ala Arg Gln Val Phe Pro Val Thr Asp 275 280 285

Val Ala Asp Asp Leu Met Thr Val Glu Val Lys Ala Tyr Asp Ser Ser 290 295 300

Pro Asp Ser 305

<210> 233

<211> 684

<212> PRT

<213> Homo sapiens

<400> 233

Met Ser Ala Ile Pro Ala Glu Glu Ser Asp Gln Leu Leu Ile Arg Pro 1 5 10 15

Leu Gly Ala Gly Gln Glu Val Gly Arg Ser Cys Ile Ile Leu Glu Phe 20 25 30

Lys Gly Arg Lys Ile Met Leu Asp Cys Gly Ile His Pro Gly Leu Glu 35 40 45

Gly Met Asp Ala Leu Pro Tyr Ile Asp Leu Ile Asp Pro Ala Glu Ile 50 55 60

Asp Leu Leu Leu Ile Ser His Phe His Leu Asp His Cys Gly Ala Leu 65 70 75 80

Pro Trp Phe Leu Gln Lys Thr Ser Phe Lys Gly Arg Thr Phe Met Thr 85 90 95

His Ala Thr Lys Ala Ile Tyr Arg Trp Leu Leu Ser Asp Tyr Val Lys 105 Val Ser Asn Ile Ser Ala Asp Asp Met Leu Tyr Thr Glu Thr Asp Leu 120 Glu Glu Ser Met Asp Lys Ile Glu Thr Ile Asn Phe His Glu Val Lys Glu Val Ala Gly Ile Lys Phe Trp Cys Tyr His Ala Gly His Val Leu Gly Ala Ala Met Phe Met Ile Glu Ile Ala Gly Val Lys Leu Leu Tyr Thr Gly Asp Phe Ser Arg Gln Glu Asp Arg His Leu Met Ala Ala Glu 185 Ile Pro Asn Ile Lys Pro Asp Ile Leu Ile Ile Glu Ser Thr Tyr Gly 195 Thr His Ile His Glu Lys Arg Glu Glu Arg Glu Ala Arg Phe Cys Asn Thr Val His Asp Ile Val Asn Arg Gly Gly Arg Gly Leu Ile Pro Val 230 Phe Ala Leu Gly Arg Ala Gln Glu Leu Leu Leu Ile Leu Asp Glu Tyr 250 Trp Gln Asn His Pro Glu Leu His Asp Ile Pro Ile Tyr Tyr Ala Ser Ser Leu Ala Lys Lys Cys Met Ala Val Tyr Gln Thr Tyr Val Asn Ala Met Asn Asp Lys Ile Arg Lys Gln Ile Asn Ile Asn Asn Pro Phe Val Phe Lys His Ile Ser Asn Leu Lys Ser Met Asp His Phe Asp Asp Ile 315 Gly Pro Ser Val Val Met Ala Ser Pro Gly Met Met Gln Ser Gly Leu Ser Arg Glu Leu Phe Glu Ser Trp Cys Thr Asp Lys Arg Asn Gly Val Ile Ile Ala Gly Tyr Cys Val Glu Gly Thr Leu Ala Lys His Ile Met 360 Ser Glu Pro Glu Glu Ile Thr Thr Met Ser Gly Gln Lys Leu Pro Leu Lys Met Ser Val Asp Tyr Ile Ser Phe Ser Ala His Thr Asp Tyr Gln 390 Gln Thr Ser Glu Phe Ile Arg Ala Leu Lys Pro Pro His Val Ile Leu 405 410 415 Val His Gly Glu Gln Asn Glu Met Ala Arg Leu Lys Ala Ala Leu Ile

Arg Glu Tyr Glu Asp Asn Asp Glu Val His Ile Glu Val His Asn Pro 440 Arg Asn Thr Glu Ala Val Thr Leu Asn Phe Arg Gly Glu Lys Leu Ala 460 Lys Val Met Gly Phe Leu Ala Asp Lys Lys Pro Glu Gln Gly Gln Arg Val Ser Gly Ile Leu Val Lys Arg Asn Phe Asn Tyr His Ile Leu Ser Pro Cys Asp Leu Ser Asn Tyr Thr Asp Leu Ala Met Ser Thr Val Lys 505 Gln Thr Gln Ala Ile Pro Tyr Thr Gly Pro Phe Asn Leu Leu Cys Tyr Gln Leu Gln Lys Leu Thr Gly Asp Val Glu Glu Leu Glu Ile Gln Glu Lys Pro Ala Leu Lys Val Phe Lys Asn Ile Thr Val Ile Gln Glu Pro 555 Gly Met Val Val Leu Glu Trp Leu Ala Asn Pro Ser Asn Asp Met Tyr 565 Ala Asp Thr Val Thr Val Ile Leu Glu Val Gln Ser Asn Pro Lys 585 Ile Arg Lys Gly Ala Val Gln Lys Val Ser Lys Lys Leu Glu Met His Val Tyr Ser Lys Arg Leu Glu Ile Met Leu Gln Asp Ile Phe Gly Glu 615 Asp Cys Val Ser Val Lys Asp Asp Ser Ile Leu Ser Val Thr Val Asp Gly Lys Thr Ala Asn Leu Asn Leu Glu Thr Arg Thr Val Glu Cys Glu Glu Gly Ser Glu Asp Asp Glu Ser Leu Arg Glu Met Val Glu Leu Ala Ala Gln Arg Leu Tyr Glu Ala Leu Thr Pro Val His

<210> 234 <211> 693

<212> PRT <213> Arabidopsis thaliana

Arg Asp Gly Asp Gln Leu Ile Val Thr Pro Leu Gly Ala Gly Ser Glu
20 25 30

Val Gly Arg Ser Cys Val Tyr Met Ser Phe Arg Gly Lys Asn Ile Leu 35 45 Phe Asp Cys Gly Ile His Pro Ala Tyr Ser Gly Met Ala Ala Leu Pro Tyr Phe Asp Glu Ile Asp Pro Ser Ser Ile Asp Val Leu Leu Ile Thr His Phe His Ile Asp His Ala Ala Ser Leu Pro Tyr Phe Leu Glu Lys Thr Thr Phe Asn Gly Arg Val Phe Met Thr His Ala Thr Lys Ala Ile Tyr Lys Leu Leu Thr Asp Tyr Val Lys Val Ser Lys Val Ser Val Glu Asp Met Leu Phe Asp Glu Gln Asp Ile Asn Lys Ser Met Asp Lys Ile Glu Val Ile Asp Phe His Gln Thr Val Glu Val Asn Gly Ile Lys 150 Phe Trp Cys Tyr Thr Ala Gly His Val Leu Gly Ala Ala Met Phe Met Val Asp Ile Ala Gly Val Arg Ile Leu Tyr Thr Gly Asp Tyr Ser Arg Glu Glu Asp Arg His Leu Arg Ala Ala Glu Leu Pro Gln Phe Ser Pro 200 Asp Ile Cys Ile Ile Glu Ser Thr Ser Gly Val Gln Leu His Gln Ser Arg His Ile Arg Glu Lys Arg Phe Thr Asp Val Ile His Ser Thr Val 230 Ala Gln Gly Gly Arg Val Leu Ile Pro Ala Phe Ala Leu Gly Arg Ala 250 Gln Glu Léu Leu Ile Leu Asp Glu Tyr Trp Ala Asn His Pro Asp Leu His Asn Ile Pro Ile Tyr Tyr Ala Ser Pro Leu Ala Lys Lys Cys 280 Met Ala Val Tyr Gln Thr Tyr Ile Leu Ser Met Asn Asp Arg Ile Arg 295 Asn Gln Phe Ala Asn Ser Asn Pro Phe Val Phe Lys His Ile Ser Pro 310 Leu Asn Ser Ile Asp Asp Phe Asn Asp Val Gly Pro Ser Val Val Met Ala Thr Pro Gly Gly Leu Gln Ser Gly Leu Ser Arg Gln Leu Phe Asp 345 Ser Trp Cys Ser Asp Lys Lys Asn Ala Cys Ile Ile Pro Gly Tyr Met Val Glu Gly Thr Leu Ala Lys Thr Ile Ile Asn Glu Pro Lys Glu Val 375

Pro Leu Ser Ala Ser

690

Thr Leu Met Asn Gly Leu Thr Ala Pro Leu Asn Met Gln Val His Tyr 395 Ile Ser Phe Ser Ala His Ala Asp Tyr Ala Gln Thr Ser Thr Phe Leu Lys Glu Leu Met Pro Pro Asn Ile Ile Leu Val His Gly Glu Ala Asn 425 Glu Met Met Arg Leu Lys Gln Lys Leu Leu Thr Glu Phe Pro Asp Gly Asn Thr Lys Ile Met Thr Pro Lys Asn Cys Glu Ser Val Glu Met Tyr Phe Asn Ser Glu Lys Leu Ala Lys Thr Ile Gly Arg Leu Ala Glu Lys Thr Pro Asp Val Gly Asp Thr Val Ser Gly Ile Leu Val Lys Lys Gly 490 Phe Thr Tyr Gln Ile Met Ala Pro Asp Glu Leu His Val Phe Ser Gln Leu Ser Thr Ala Thr Val Thr Gln Arg Ile Thr Ile Pro Phe Val Gly Ala Phe Gly Val Ile Lys His Arg Leu Glu Lys Ile Phe Glu Ser Val 535 Glu Phe Ser Thr Asp Glu Glu Ser Gly Leu Pro Ala Leu Lys Val His 550 Glu Arg Val Thr Val Lys Gln Glu Ser Glu Lys His Ile Ser Leu Gln 570 Trp Ser Ser Asp Pro Ile Ser Asp Met Val Ser Asp Ser Ile Val Ala 580 Leu Ile Leu Asn Ile Ser Arg Glu Val Pro Lys Ile Val Met Glu Glu 600 Glu Asp Ala Val Lys Ser Glu Glu Glu Asn Gly Lys Lys Val Glu Lys 610 Val Ile Tyr Ala Leu Leu Val Ser Leu Phe Gly Asp Val Lys Leu Gly 635 Glu Asn Gly Lys Leu Val Ile Arg Val Asp Gly Asn Val Ala Gln Leu 650 Asp Lys Glu Ser Gly Glu Val Glu Ser Glu His Ser Gly Leu Lys Glu Arg Val Arg Val Ala Phe Glu Arg Ile Gln Ser Ala Val Lys Pro Ile 680

<210> 235

<211> 779 <212> PRT

<213> Saccharomyces cerevisiae

<400> 235

Met Glu Arg Thr Asn Thr Thr Thr Phe Lys Phe Phe Ser Leu Gly Gly
1 5 10 15

Ser Asn Glu Val Gly Arg Ser Cys His Ile Leu Gln Tyr Lys Gly Lys 20 25 30

Thr Val Met Leu Asp Ala Gly Ile His Pro Ala Tyr Gln Gly Leu Ala 35 40 45

Ser Leu Pro Phe Tyr Asp Glu Phe Asp Leu Ser Lys Val Asp Ile Leu 50 55 60

Leu Ile Ser His Phe His Leu Asp His Ala Ala Ser Leu Pro Tyr Val 65 70 75 80

Met Gln Arg Thr Asn Phe Gln Gly Arg Val Phe Met Thr His Pro Thr 85 90 95

Lys Ala Ile Tyr Arg Trp Leu Leu Arg Asp Phe Val Arg Val Thr Ser 100 105 110

Ile Gly Ser Ser Ser Ser Met Gly Thr Lys Asp Glu Gly Leu Phe 115 120 125

Ser Asp Glu Asp Leu Val Asp Ser Phe Asp Lys Ile Glu Thr Val Asp 130 135 140

Tyr His Ser Thr Val Asp Val Asn Gly Ile Lys Phe Thr Ala Phe His 145 150 155 160

Ala Gly His Val Leu Gly Ala Ala Met Phe Gln Ile Glu Ile Ala Gly 165 170 175

Leu Arg Val Leu Phe Thr Gly Asp Tyr Ser Arg Glu Val Asp Arg His 180 185 190

Leu Asn Ser Ala Glu Val Pro Pro Leu Ser Ser Asn Val Leu Ile Val 195 200 205

Glu Ser Thr Phe Gly Thr Ala Thr His Glu Pro Arg Leu Asn Arg Glu 210 215 220

Arg Lys Leu Thr Gln Leu Ile His Ser Thr Val Met Arg Gly Gly Arg 225 230 235 240

Val Leu Leu Pro Val Phe Ala Leu Gly Arg Ala Gln Glu Ile Met Leu 245 250 255

Ile Leu Asp Glu Tyr Trp Ser Gln His Ala Asp Glu Leu Gly Gly Gly 260 265 270

Gln Val Pro Ile Phe Tyr Ala Ser Asn Leu Ala Lys Lys Cys Met Ser 275 280 285

Val Phe Gln Thr Tyr Val Asn Met Met Asn Asp Asp Ile Arg Lys Lys 290 295 300

Phe Arg Asp Ser Gln Thr Asn Pro Phe Ile Phe Lys Asn Ile Ser Tyr 315 310 Leu Arg Asn Leu Glu Asp Phe Gln Asp Phe Gly Pro Ser Val Met Leu 330 Ala Ser Pro Gly Met Leu Gln Ser Gly Leu Ser Arg Asp Leu Leu Glu 345 Arg Trp Cys Pro Glu Asp Lys Asn Leu Val Leu Ile Thr Gly Tyr Ser Ile Glu Gly Thr Met Ala Lys Phe Ile Met Leu Glu Pro Asp Thr Ile 375 Pro Ser Ile Asn Asn Pro Glu Ile Thr Ile Pro Arg Arg Cys Gln Val Glu Glu Ile Ser Phe Ala Ala His Val Asp Phe Gln Glu Asn Leu Glu Phe Ile Glu Lys Ile Ser Ala Pro Asn Ile Ile Leu Val His Gly Glu 425 Ala Asn Pro Met Gly Arg Leu Lys Ser Ala Leu Leu Ser Asn Phe Ala Ser Leu Lys Gly Thr Asp Asn Glu Val His Val Phe Asn Pro Arg Asn 455 Cys Val Glu Val Asp Leu Glu Phe Gln Gly Val Lys Val Ala Lys Ala Val Gly Asn Ile Val Asn Glu Ile Tyr Lys Glu Glu Asn Val Glu Ile 490 Lys Glu Glu Ile Ala Ala Lys Ile Glu Pro Ile Lys Glu Glu Asn Glu Asp Asn Leu Asp Ser Gln Ala Glu Lys Gly Leu Val Asp Glu Glu Glu 520 His Lys Asp Ile Val Val Ser Gly Ile Leu Val Ser Asp Asp Lys Asn Phe Glu Leu Asp Phe Leu Ser Leu Ser Asp Leu Arg Glu His His Pro 550 Asp Leu Ser Thr Thr Ile Leu Arg Glu Arg Gln Ser Val Arg Val Asn 570 Cys Lys Lys Glu Leu Ile Tyr Trp His Ile Leu Gln Met Phe Gly Glu 580 Ala Glu Val Leu Gln Asp Asp Asp Arg Val Thr Asn Gln Glu Pro Lys Val Lys Glu Glu Ser Lys Asp Asn Leu Thr Asn Thr Gly Lys Leu Ile Leu Gln Ile Met Gly Asp Ile Lys Leu Thr Ile Val Asn Thr Leu Ala 630

Val Val Glu Trp Thr Gln Asp Leu Met Asn Asp Thr Val Ala Asp Ser 645 650 655

Ile Ile Ala Ile Leu Met Asn Val Asp Ser Ala Pro Ala Ser Val Lys 660 665 670

Leu Ser Ser His Ser Cys Asp Asp His Asp His Asn Asn Val Gln Ser 675 680 685

Asn Ala Gln Gly Lys Ile Asp Glu Val Glu Arg Val Lys Gln Ile Ser 690 695 700

Arg Leu Phe Lys Glu Gln Phe Gly Asp Cys Phe Thr Leu Phe Leu Asn 705 710 715 720

Lys Asp Glu Tyr Ala Ser Asn Lys Glu Glu Thr Ile Thr Gly Val Val 725 730 735

Thr Ile Gly Lys Ser Thr Ala Lys Ile Asp Phe Asn Asn Met Lys Ile 740 745 750

Leu Glu Cys Asn Ser Asn Pro Leu Lys Gly Arg Val Glu Ser Leu Leu 755 760 765

Asn Ile Gly Gly Asn Leu Val Thr Pro Leu Cys 770 775

<210> 236

<211> 554

<212> PRT

<213> Synechocystis sp.

<400> 236 Met Thr Phe Ser Val Pro Thr Gln Gly Lys Ala Phe Ala Asn Ile Ser

Phe Leu Pro Tyr Gly Val Gly Pro Arg Asp Gly Gly Ile Cys Leu Glu 20 25 30

Leu His Leu Gly Pro Tyr Arg Ile Leu Leu Asp Cys Gly Leu Glu Asp

Leu Thr Pro Leu Leu Ala Ala Asp Pro Gly Thr Val Asp Leu Val Phe
50 55 60

Cys Ser His Ala His Arg Asp His Gly Leu Gly Leu Trp Gln Phe His 65 70 75 80

Gln Gln Phe Pro His Ile Pro Ile Leu Ala Ser Glu Val Thr Gln Arg 85 90 95

Leu Leu Pro Leu Asn Trp Pro Asp Glu Phe Val Pro Pro Phe Cys Arg 100 105 110

Val Leu Pro Trp Arg Ser Pro Gln Glu Val Leu Pro Gly Leu Thr Val 115 120 125

Glu Leu Leu Pro Ala Gly His Leu Pro Gly Ala Ala Leu Ile Leu Leu 130 135 140

Glu Tyr His Asn Gly Asp Arg Leu Tyr Arg Val Ile Tyr Thr Gly Asp 145 150 155

Tyr Cys Leu Ser His Leu Gln Leu Val Asp Gly Leu Ala Leu Thr Pro 165 Leu Arg Gly Leu Lys Pro Asp Val Leu Ile Leu Glu Gly His Tyr Gly Asn Arg Arg Leu Pro His Arg Arg Gln Gln Glu Lys Gln Phe Ile Gln Ala Ile Glu Thr Val Leu Ala Lys Gly Arg Asn Ile Leu Leu Pro Val Pro Pro Leu Gly Leu Ala Gln Glu Ile Leu Lys Leu Leu Arg Thr His His Gln Phe Thr Gly Arg Gln Val Asn Leu Trp Ala Gly Glu Ser Val 250 Ala Arg Gly Cys Asp Ala Tyr Gln Gly Ile Ile Asp His Leu Pro Asp Asn Val Arg Asn Phe Ala Gln His Gln Pro Leu Phe Trp Asp Asp Lys Val Tyr Pro His Leu Arg Pro Leu Thr Asp Asp Gln Gly Glu Leu Ser Leu Ser Ala Pro Ser Ile Val Ile Thr Thr Thr Trp Pro Ala Phe Trp 310 Pro Ser Pro Ala Ala Leu Pro Gly Leu Trp Thr Val Phe Met Pro Gln Leu Leu Thr Leu Pro Ser Cys Leu Val Asn Phe Ala Trp Gln Asp Leu 345 Glu Glu Phe Pro Lys Tyr Glu Leu Glu Asp Tyr Leu Leu Ala Asp His Ser Asp Gly Arg Asn Thr Thr Gln Leu Ile His Asn Leu Arg Pro Gln His Leu Val Phe Val His Gly Gln Pro Ser Asp Ile Glu Asp Leu Thr 395 Ser Leu Glu Glu Leu Gln Ser Arg Tyr Gln Leu His Ser Pro Ala Ala 405 Gly Asn Ala Val Ala Leu Pro Ile Gly Asp Arg Phe Val Gln Pro Thr 425 Pro Pro Pro Gln Ile Tyr Glu Gly Glu Ile His Glu Leu Glu Pro Asn Lys Gln Ile His His Leu Gly Glu Val Val Ile His Leu Asp Gly 455 Gln Ile Leu Glu Asn Ser Arg Trp Gly Lys Phe Gly Glu Thr Gly Ile 470 Val Gln Ala Arg Trp Gln Gly Glu Glu Leu Val Leu Arg Gly Ile Ser

Gln Arg Glu Leu Leu Lys Gln Asn Gln Ser Ser Lys Arg Pro Val Asp 500 505 510

Phe Asp Cys Cys Ala Asn Cys Arg His Phe Gln His Tyr His Cys Arg 515 520 525

Asn Pro Val Ser Pro Leu Met Gly Leu Glu Val Arg Ala Asp Gly His 530 535 540

Cys Pro Val Phe Glu Ser Val Ala Ser Ser 545 550

<210> 237

<211> 636

<212> PRT

<213> Methanobacterium thermoautotrophicum

<400> 237 .

Met Val Ser Glu Met Leu Glu Glu Ile Lys Arg Thr Ile Met Gln Arg 1 5 10 15

Leu Pro Glu Arg Val Gln Val Ala Lys Val Glu Phe Glu Gly Pro Glu 20 25 30

Val Val Ile Tyr Thr Lys Asn Pro Glu Ile Ile Thr Glu Asn Gly Asn 35 40 45

Leu Ile Arg Asp Ile Ala Lys Asp Ile Arg Lys Arg Ile Ile Ile Arg 50 55 60

Ser Asp Arg Ser Val Leu Met Asp Pro Glu Lys Ala Ile Arg Lys Ile 65 70 75 80

His Glu Ile Val Pro Glu Glu Ala Lys Ile Thr Asn Ile Ser Phe Asp 85 90 95

Asp Val Thr Cys Glu Val Ile Ile Glu Ala Arg Lys Pro Gly Leu Val 100 105 110

Ile Gly Lys Tyr Gly Ser Thr Ser Arg Glu Ile Val Lys Asn Thr Gly 115 120 125

Trp Ala Pro Lys Ile Leu Arg Thr Pro Pro Ile Ser Ser Glu Ile Ile 130 135 140

Glu Arg Ile Arg Arg Thr Leu Arg Lys Asn Ser Lys Glu Arg Lys Lys 145 150 155 160

Ile Leu Gln Gln Leu Gly Asn Arg Ile His Gln Lys Pro Lys Tyr Asp 165 170 175

Asn Asp Trp Ala Arg Leu Thr Ala Met Gly Gly Phe Arg Glu Val Gly 180 185 190

Arg Ser Cys Leu Tyr Leu Gln Thr Pro Asn Ser Arg Val Leu Leu Asp 195 200 205

Cys Gly Val Asn Val Ala Gly Gly Asp Asp Lys Asn Ser Tyr Pro Tyr 210 215 220

Leu Asn Val Pro Glu Phe Thr Leu Asp Ser Leu Asp Ala Val Ile Ile 225 230 235 240

Thr His Ala His Leu Asp His Ser Gly Phe Leu Pro Tyr Leu Tyr His Tyr Gly Tyr Asp Gly Pro Val Tyr Cys Thr Ala Pro Thr Arg Asp Leu Met Thr Leu Leu Gln Leu Asp His Ile Asp Ile Ala His Arg Glu Asp Glu Pro Leu Pro Phe Asn Val Lys His Val Lys Lys Ser Val Lys His Thr Ile Thr Leu Asp Tyr Gly Glu Val Thr Asp Ile Ala Pro Asp Ile Arg Leu Thr Leu His Asn Ala Gly His Ile Leu Gly Ser Ala Met Ala 325 His Leu His Ile Gly Asp Gly Gln His Asn Met Val Tyr Thr Gly Asp Phe Lys Tyr Glu Gln Ser Arg Leu Leu Glu Ala Ala Ala Asn Arg Phe Pro Arg Ile Glu Thr Leu Val Met Glu Ser Thr Tyr Gly Gly His Glu 375 Asp Val Gln Pro Ser Arg Asn Arg Ala Glu Lys Glu Leu Val Lys Thr 390 Ile Tyr Ser Thr Leu Arg Arg Gly Gly Lys Ile Leu Ile Pro Val Phe Ala Val Gly Arg Ala Gln Glu Leu Met Ile Val Leu Glu Glu Tyr Ile 425 Arg Thr Gly Ile Ile Asp Glu Val Pro Val Tyr Ile Asp Gly Met Ile Trp Glu Ala Asn Ala Ile His Thr Ala Arg Pro Glu Tyr Leu Ser Lys 460 Asp Leu Arg Asp Gln Ile Phe His Met Gly His Asn Pro Phe Ile Ser 470 Asp Ile Phe His Lys Val Asn Gly Met Asp Glu Arg Arg Glu Ile Val Glu Gly Glu Pro Ser Ile Ile Leu Ser Thr Ser Gly Met Leu Thr Gly 505 Gly Asn Ser Leu Glu Tyr Phe Lys Trp Leu Cys Glu Asp Pro Asp Asn Ser Leu Val Phe Val Gly Tyr Gln Ala Glu Gly Ser Leu Gly Arg Arg Ile Gln Lys Gly Trp Lys Glu Ile Pro Leu Lys Asp Glu Asp Asp Lys Met Arg Val Tyr Asn Val Arg Met Asn Ile Lys Thr Ile Glu Gly Phe

Ser Gly His Ser Asp Arg Arg Gln Leu Met Glu Tyr Val Lys Arg Ile 580 585 590

Ser Pro Lys Pro Glu Lys Ile Leu Leu Cys His Gly Asp Asn Tyr Lys 595 600 605

Thr Leu Asp Leu Ala Ser Ser Ile Tyr Arg Thr Tyr Arg Ile Glu Thr 610 615 620

Lys Thr Pro Leu Asn Leu Glu Thr Val Arg Ile Gln 625 635

<210> 238

<211> 1040

<212> PRT

<213> Homo sapiens

<400> 238

Met Leu Glu Asp Ile Ser Glu Glu Asp Ile Trp Glu Tyr Lys Ser Lys

1 10 15

Arg Lys Pro Lys Arg Val Asp Pro Asn Asn Gly Ser Lys Asn Ile Leu 20 · 25 30

Lys Ser Val Glu Lys Ala Thr Asp Gly Lys Tyr Gln Ser Lys Arg Ser 35 40 45

Arg Asn Arg Lys Arg Ala Ala Glu Ala Lys Glu Val Lys Asp His Glu
50 55 60

Val Pro Leu Gly Asn Ala Gly Cys Gln Thr Ser Val Ala Ser Ser Gln 65 70 75 80

Asn Ser Ser Cys Gly Asp Gly Ile Gln Gln Thr Gln Asp Lys Glu Thr 85 90 95

Thr Pro Gly Lys Leu Cys Arg Thr Gln Lys Ser Gln His Val Ser Pro 100 105 110

Lys Ile Arg Pro Val Tyr Asp Gly Tyr Cys Pro Asn Cys Gln Met Pro 115 120 125

Phe Ser Ser Leu Ile Gly Gln Thr Pro Arg Trp His Val Phe Glu Cys 130 140

Leu Asp Ser Pro Pro Arg Ser Glu Thr Glu Cys Pro Asp Gly Leu Leu 145 150 155 160

Cys Thr Ser Thr Ile Pro Phe His Tyr Lys Arg Tyr Thr His Phe Leu

Leu Ala Gln Ser Arg Ala Gly Asp His Pro Phe Ser Ser Pro Ser Pro 180 185 190

Ala Ser Gly Gly Ser Phe Ser Glu Thr Lys Ser Gly Val Leu Cys Ser 195 200 205

Leu Glu Glu Arg Trp Ser Ser Tyr Gln Asn Gln Thr Asp Asn Ser Val 210 225 220

Ser Asn Asp Pro Leu Leu Met Thr Gln Tyr Phe Lys Lys Ser Pro Ser 225 230 235

Leu Thr Glu Ala Ser Glu Lys Ile Ser Thr His Ile Gln Thr Ser Gln Gln Ala Leu Gln Phe Thr Asp Phe Val Glu Asn Asp Lys Leu Val Gly 265 Val Ala Leu Arg Leu Ala Asn Asn Ser Glu His Ile Asn Leu Pro Leu Pro Glu Asn Asp Phe Ser Asp Cys Glu Ile Ser Tyr Ser Pro Leu Gln Ser Asp Glu Asp Thr His Asp Ile Asp Glu Lys Pro Asp Asp Ser Gln Glu Gln Leu Phe Phe Thr Glu Ser Ser Lys Asp Gly Ser Leu Glu Glu Asp Asp Asp Ser Cys Gly Phe Phe Lys Lys Arg His Gly Pro Leu Leu Lys Asp Gln Asp Glu Ser Cys Pro Lys Val Asn Ser Phe Leu Thr Arg 360 Asp Lys Tyr Asp Glu Gly Leu Tyr Arg Phe Asn Ser Leu Asn Asp Leu Ser Gln Pro Ile Ser Gln Asn Asn Glu Ser Thr Leu Pro Tyr Asp Leu 395 390 Ala Cys Thr Gly Gly Asp Phe Val Leu Phe Pro Pro Ala Leu Ala Gly 410 Lys Leu Ala Ala Ser Val His Gln Ala Thr Lys Ala Lys Pro Asp Glu 425 420 Pro Glu Phe His Ser Ala Gln Ser Asn Lys Gln Lys Gln Val Ile Glu Glu Ser Ser Val Tyr Asn Gln Val Ser Leu Pro Leu Val Lys Ser Leu Met Leu Lys Pro Phe Glu Ser Gln Val Glu Gly Tyr Leu Ser Ser Gln 470 Pro Thr Gln Asn Thr Ile Arg Lys Leu Ser Ser Glu Asn Leu Asn Ala Lys Asn Asn Thr Asn Ser Ala Cys Phe Cys Arg Lys Ala Leu Glu Gly 505 Val Pro Val Gly Lys Ala Thr Ile Leu Asn Thr Glu Asn Leu Ser Ser 520 Thr Pro Ala Pro Lys Tyr Leu Lys Ile Leu Pro Ser Gly Leu Lys Tyr Asn Ala Arg His Pro Ser Thr Lys Val Met Lys Gln Met Asp Ile Gly 550 Val Tyr Phe Gly Leu Pro Pro Lys Arg Lys Glu Glu Lys Leu Leu Gly

Glu	Ser	Ala	Leu 580	Glu	Gly	Ile	Asn	Leu 585	Asn	Pro	Val	Pro	Ser 590	Pro	Asn
Gln	Lys	Arg 595	Ser	Ser	Gln	Cys	Lys 600	Arg	Lys	Ala	Glu	Lys 605	Ser	Leu	Ser
Asp	Leu 610	Glu	Phe	Asp	Ala	Ser 615	Thr	Leu	His	Glu	Ser 620	Gln	Leu	Ser	Val
Glu 625	Leu	Ser	Ser	Glu	Arg 630	Ser	Gln _.	Arg	Gln	Lys 635	Lys	Arg	Cys	Arg	Lys 640
Ser	Asn	Ser	Leu	Gln 645	Glu	Gly	Ala	Cys	Gln 650	Lys	Arg	Ser	Asp	His 655	Leu
Ile	Asn	Thr	Glu 660	Ser	Glu	Ala	Val	Asn 665	Leu	Ser	Lys	Val	Lys 670	Val	Phe
Thr	Lys	Ser 675	Ala	His	Gly	Gly	Leu 680	Gln	Arg	Gly	Asn	Lys 685	Lys	Ile	Pro
Glu	Ser 690	Ser	Asn	Val	Gly	Gly 695	Ser	Arg	Lys	Lys	Thr 700	Суѕ	Pro	Phe	Tyr
Lys 705	Lys	Ile	Pro	Gly	Thr 710	Gly	Phe	Thr	Val	Asp 715	Ala	Phe	Gln	Tyr	Gly 720
Val	Val	Glu	Gly	Cys 725	Thr	Ala	Tyr	Phe	Leu 730	Thr	His	Phe	His	Ser 735	Asp
His	Tyr	Ala	Gly 740	Leu	Ser	Lys	His	Phe 745	Thr	Phe	Pro	Val	Tyr 750	Cys	Ser
Glu	Ile	Thr 755	Gly	Asn	Leu	Leu	Lys 760	Asn	Lys	Leu	His	Val 765	Gln	Glu	Gln
Tyr	Ile 770	His	Pro	Leu	Pro	Leu 775	Asp	Thr	Glu	Cys	Ile 780	Val	Asn	Gly	Val
Lys 785	Val	Val	Leu	Leu	Asp 790	Ala	Asn	His	Cys	Pro 795	Gly	Ala	Val	Met	Ile 800
Leu	Phe	Tyr	Leu	Pro 805	Asn	Gly	Thr	Val	Ile 810	Leu	His	Thr	Gly	Asp 815	Phe
Arg	Ala	Asp	Pro 820	Ser	Met	Glu	Arg	Ser 825	Leu	Leu	Ala	Asp	Gln 830	Lys	Val
His	Met	Leu 835	Tyr	Leu	Asp	Thr	Thr 840	Tyr	Cys	Ser	Pro	Glu 845	Туг	Thr	Phe
Pro	Ser 850		Gln	Glu	Val	Ile 855	Arg	Phe	Ala	Ile	860	Thr	Ala	Phe	e Glu
Ala 865		Thr	Leu	Asn	Pro 870		Ala	Leu	Val	Val 875	. Cys	Gly	Th:	Туг	Ser 880
Ile	Gly	Lys	Glu	Lys 885		Phe	Leu	Ala	11e	e Ala	a Asp	Val	. Le	3 Gly 895	y Ser
Lys	Val	Gly	Met 900		Gln	Glu	Lys	Tyr 905	Lys	s Thr	r Lev	ı Glr	n Cy: 91	s Lev	ı Asn

Ile Pro Glu Ile Asn Ser Leu Ile Thr Thr Asp Met Cys Ser Ser Leu 920 Val His Leu Leu Pro Met Met Gln Ile Asn Phe Lys Gly Leu Gln Ser 935 930 His Leu Lys Lys Cys Gly Gly Lys Tyr Asn Gln Ile Leu Ala Phe Arg Pro Thr Gly Trp Thr His Ser Asn Lys Phe Thr Arg Ile Ala Asp Val 970 Ile Pro Gln Thr Lys Gly Asn Ile Ser Ile Tyr Gly Ile Pro Tyr Ser Glu His Ser Ser Tyr Leu Glu Met Lys Arg Phe Val Gln Trp Leu Lys 1000 Pro Gln Lys Ile Ile Pro Thr Val Asn Val Gly Thr Trp Lys Ser Arg 1015 Ser Thr Met Glu Lys Tyr Phe Arg Glu Trp Lys Leu Glu Ala Gly Tyr 1030 1025

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<213> Arabidopsis thaliana

<400> 239

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Ser Ser Gln Leu Ser Ile Arg Lys Pro Leu His Pro Thr Asn Ala Asn 20 25 30

Asn Ile Ser His Arg Pro Pro Asn Lys Lys Pro Arg Leu Cys Arg Tyr 35 40 45

Pro Gly Lys Glu Asn Val Thr Pro Pro Pro Ser Pro Asp Pro Asp Leu
50 55 60

Phe Cys Ser Ser Ser Thr Pro His Cys Ile Leu Asp Cys Ile Pro Ser 65 70 75

Ser Val Asp Cys Ser Leu Gly Asp Phe Asn Gly Pro Ile Ser Ser Leu
85 90 95

Gly Glu Glu Asp Lys Glu Asp Lys Asp Asp Cys Ile Lys Val Asn Arg

Glu Gly Tyr Leu Cys Asn Ser Met Glu Ala Arg Leu Leu Lys Ser Arg 115 120 125

Ile Cys Leu Gly Phe Asp Ser Gly Ile His Glu Asp Asp Glu Gly Phe 130 135 140

Val Glu Ser Asn Ser Glu Leu Asp Val Leu Ile Asn Leu Cys Ser Glu 145 150 155 160

Ser	Glu	Gly	Arg	Ser 165	Gly	Glu	Phe	Ser	Leu 170	Gly	Lys	Asp	Asp	Ser 175	Ile
Gln	Cys	Pro	Leu 180	Cys	Ser	Met	Asp	Ile 185	Ser	Ser	Leu	Ser	Glu 190	Glu	Gln
Arg	Gln	Val 195	His	Ser	Asn	Thr	Cys 200	Leu	Asp	Lys	Ser	Tyr 205	Asn	Gln	Pro
Ser	Glu 210	Gln	Asp	Ser	Leu	Arg 215	Lys	Cys	Glu	Asn	Leu 220	Ser	Ser	Leu	Ile
Lys 225	Glu	Ser	Ile	Asp	Asp 230	Pro	Val	Gln	Leu	Pro 235	Gln	Leu	Val	Thr	Asp 240
Leu	Ser	Pro	Val	Leu 245	Lys	Trp	Leu	Arg	Ser 250	Leu	Gly	Leu	Ala	Lys 255	Tyr
Glu	Asp	Val	Phe 260	Ile	Arg	Glu	Glu	Ile 265	Asp	Trp	Asp	Thr	Leu 270	Gln	Ser
Leu	Thr	Glu 275	Glu	Asp	Leu	Leu	Ser 280	Ile	Gly	Ile	Thr	Ser 285	Leu	Gly	Pro
Arg	Lys 290	Lys	Ile	Val	Asn	Ala 295	Leu	Ser	Gly	Val	Arg 300	Asp	Pro	Phe	Ala
Ser 305	Ser	Ala	Glu	Val	Gln 310	Ala	Gln	Ser	His	Cys 315	Thr	Ser	Gly	His	Val 320
Thr	Glu	Arg	Gln	Arg 325	Asp	Lys	Ser	Thr	Thr 330	Arg	Lys	Ala	Ser	Glu 335	Pro
Lys	Lys	Pro	Thr 340	Ala	Asn	Lys	Leu	Ile 345	Thr	Glu	Phe	Phe	Pro 350	Gly	Gln
Ala	Thr	Glu 355	Gly	Thr	Lys	Ile	Arg 360	Thr	Ala	Pro	Lys	Pro 365	Val	Ala	Glu
Lys	Ser 370	Pro	Ser	Asp	Ser	Ser 375	Ser	Arg	Arg	Ala	Val 380	Arg	Arg	Asn	Gly
Asn 385	Asn	Gly	Lys	Ser	Lys 390	Val	Ile	Pro	His	Trp 395	Asn	Cys	Ile	Pro	Gly 400
Thr	Pro	Phe	Arg	Val 405	Asp	Ala	Phe	Lys	Tyr 410	Leu	Thr	Arg	Asp	Cys 415	Cys
His	Trp	Phe	Leu 420	Thr	His	Phe	His	Leu 425		His	Tyr	Gln	Gly 430	Leu	Thr
Lys	Ser	Phe 435		His	Gly	Lys	Ile 440		Cys	Ser	Leu	Val 445	Thr	Ala	Lys
Leu	Val 450	Asn	Met	Lys	Ile	Gly 455		Pro	Trp	Glu	Arg 460	Leu	Gln	Val	Leu
Asp 465	Leu	Gly	Gln	Lys	Val 470		Ile	Ser	Gly	7 Ile 475	Asp	Val	Thr	Cys	Phe 480
Asp	Ala	Asn	His	Cys 485		Gly	Ser	Ile	Met 490	: Ile	. Leu	Phe	e Glu	495	Ala



Asn Gly Lys Ala Val Leu His Thr Gly Asp Phe Arg Tyr Ser Glu Glu 505 Met Ser Asn Trp Leu Ile Gly Ser His Ile Ser Ser Leu Ile Leu Asp 520 Thr Thr Tyr Cys Asn Pro Gln Tyr Asp Phe Pro Lys Gln Glu Ala Val Ile Gln Phe Val Val Glu Ala Ile Gln Ala Glu Ala Phe Asn Pro Lys Thr Leu Phe Leu Ile Gly Ser Tyr Thr Ile Gly Lys Glu Arg Leu Phe Leu Glu Val Ala Arg Val Leu Arg Glu Lys Ile Tyr Ile Asn Pro Ala Lys Leu Lys Leu Glu Cys Leu Gly Phe Ser Lys Asp Asp Ile Gln Trp Phe Thr Val Lys Glu Glu Glu Ser His Ile His Val Val Pro Leu 615 Trp Thr Leu Ala Ser Phe Lys Arg Leu Lys His Val Ala Asn Arg Tyr Thr Asn Arg Tyr Ser Leu Ile Val Ala Phe Ser Pro Thr Gly Trp Thr Ser Gly Lys Thr Lys Lys Ser Pro Gly Arg Arg Leu Gln Gln Gly Thr Ile Ile Arg Tyr Glu Val Pro Tyr Ser Glu His Ser Ser Phe Thr Glu Leu Lys Glu Phe Val Gln Lys Val Ser Pro Glu Val Ile Ile Pro Ser Val Asn Asn Asp Gly Pro Asp Ser Ala Ala Ala Met Val Ser Leu 710

<210> 240

Leu Val Thr

<211> 661

<212> PRT

<213> Saccharomyces cerevisiae

<400> 240

Met Ser Arg Lys Ser Ile Val Gln Ile Arg Arg Ser Glu Val Lys Arg

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Lys Arg Ser Ser Thr Ala Ser Ser Thr Ser Glu Gly Lys Thr Leu His
20 25 30

Lys Asn Thr His Thr Ser Ser Lys Arg Gln Arg Thr Leu Thr Glu Phe 35 40 45

Asn Ile Pro Thr Ser Ser Asn Leu Pro Val Arg Ser Ser Ser Tyr Ser 50 55 60





111

Phe Ser Arg Phe Ser Cys Ser Thr Ser Asn Lys Asn Thr Glu Pro Val Ile Ile Asn Asp Asp Asp His Asn Ser Ile Cys Leu Glu Asp Thr Ala Lys Val Glu Ile Thr Ile Asp Thr Asp Glu Glu Glu Leu Val Ser Leu His Asp Asn Glu Val Ser Ala Ile Glu Asn Arg Thr Glu Asp Arg Ile 120 Val Thr Glu Leu Glu Glu Gln Val Asn Val Lys Val Ser Thr Glu Val Ile Gln Cys Pro Ile Cys Leu Glu Asn Leu Ser His Leu Glu Leu Tyr Glu Arg Glu Thr His Cys Asp Thr Cys Ile Gly Ser Asp Pro Ser Asn 165 Met Gly Thr Pro Lys Lys Asn Ile Arg Ser Phe Ile Ser Asn Pro Ser Ser Pro Ala Lys Thr Lys Arg Asp Ile Ala Thr Ser Lys Lys Pro Thr Arg Val Lys Leu Val Leu Pro Ser Phe Lys Ile Ile Lys Phe Asn Asn 215 Gly His Glu Ile Val Val Asp Gly Phe Asn Tyr Lys Ala Ser Glu Thr 230 Ile Ser Gln Tyr Phe Leu Ser His Phe His Ser Asp His Tyr Ile Gly Leu Lys Lys Ser Trp Asn Asn Pro Asp Glu Asn Pro Ile Lys Lys Thr 265 Leu Tyr Cys Ser Lys Ile Thr Ala Ile Leu Val Asn Leu Lys Phe Lys Ile Pro Met Asp Glu Ile Gln Ile Leu Pro Met Asn Lys Arg Phe Trp 295 Ile Thr Asp Thr Ile Ser Val Val Thr Leu Asp Ala Asn His Cys Pro Gly Ala Ile Ile Met Leu Phe Gln Glu Phe Leu Ala Asn Ser Tyr Asp 330 325 Lys Pro Ile Arg Gln Ile Leu His Thr Gly Asp Phe Arg Ser Asn Ala 345 Lys Met Ile Glu Thr Ile Gln Lys Trp Leu Ala Glu Thr Ala Asn Glu Thr Ile Asp Gln Val Tyr Leu Asp Thr Thr Tyr Met Thr Met Gly Tyr 375 Asn Phe Pro Ser Gln His Ser Val Cys Glu Thr Val Ala Asp Phe Thr





Leu	Arg	Leu	Ile	Lys 405	His	Gly	Lys	Asn	Lys 410	Thr	Phe	Gly	Asp	Ser 415	Gln
Arg	Asn	Leu	Phe 420	His	Phe	Gln	Arg	Lys 425	Lys	Thr	Leu	Thr	Thr 430	His	Arg
Tyr	Arg	Val 435	Leu	Phe	Leu	Val	Gly 440	Thr	Tyr	Thr	Ile	Gly 445	Lys	Glu	Lys
Leu	Ala 450	Ile	Lys	Ile	Cys	Glu 455	Phe	Leu	Lys	Thr	Lys 460	Leu	Phe	Val	Met
Pro 465	Asn	Ser	Val		Phe 470	Ser	Met	Met	Leu	Thr 475	Val	Leu	Gln	Asn	Asn 480
Glu	Asn	Gln	Asn	Asp 485	Met	Trp	Asp	Glu	Ser 490	Leu	Leu	Thr	Ser	Asn 495	Leu
His	Glu	Ser	Ser 500	Val	His	Leu	Val	Pro 505	Ile	Arg	Val	Leu	Lys 510	Ser	Gln
Glu	Thr	Ile 515	Glu	Ala	Tyr	Leu	Lys 520	Ser	Leu	Lys	Glu	Leu 525	Glu	Thr	Asp
Tyr	Val 530	Lys	Asp	Ile	Glu	Asp 535	Val	Val	Gly	Phe	Ile 540	Pro	Thr	Gly	Trp
Ser 545	His	Asn	Phe	Gly	Leu 550	Lys	Tyr	Gln	Lys	Lys 555	Asn	Asp	Asp	Asp	Glu 560
.Asn	Glu	Met	Ser	Gly 565	Asn	Thr	Glu	Tyr	Cys 570	Leu	Glu	Leu	Met	Lys 575	Asn
Asp	Arg	Asp	Asn 580	Asp	Asp	Glu	Asn	Gly 585	Phe	Glu	Ile	Ser	Ser 590	Ile	Leu
Arg	Gln	Tyr 595	Lys	Lys	Tyr	Asn	Lys 600	Phe	Gln	Val	Phe	Asn 605	Val	Pro	Tyr
Ser	Glu 610	His	Ser	Ser	Phe	Asn 615	Asp	Leu	Val	Lys	Phe 620	Gly	Суѕ	Lys	Leu
Lys 625	Cys	Ser	Glu	Val	Ile 630	Pro	Thr	Val	Asn	Leu 635	Asn	Asn	Leu	Trp	Lys 640
Val	Arg	Tyr	Met	Thr 645	Asn	Trp	Phe	Gln	Cys 650	Trp	Glu	Asn	Val	Arg 655	Lys
Thr	Arg	Ala	Ala 660	Lys											